



SEQUENCE LISTING

<110> Gerlach, Robert L
MacDougall, John R
Smithson, Glenna
Millet, Isabelle
Stone, David
Gunther, Erik
Ellerman, Karen
Grosse, William M
Alsobrook II, John P
Lepley, Denise M
Burgess, Catherine E
Padigar, Muralidhara
Kekuda, Ramesh
Spytek, Kimberly A
Leach, Martin D
Shimkets, Richard A

<120> Novel Proteins and Nucleic Acids Encoding Same

B¹
<130> 21402-124

<140> 09/964,956

<141> 2001-09-26

<150> 60/235,631

<151> 2000-09-27

<150> 60/235,633

<151> 2000-09-27

<150> 60/235,808

<151> 2000-09-27

<150> 60/236,064

<151> 2000-09-27

<150> 60/236,065

<151> 2000-09-27

<150> 60/236,066

<151> 2000-09-27

<150> 60/236,135

<151> 2000-09-28

<150> 60/237,434

<151> 2000-10-03

<150> 60/238,321

<151> 2000-10-05

<150> 60/238,399

<151> 2000-10-06

<150> 60/238,396

<151> 2000-10-06

<150> 60/276,667

<151> 2001-03-16

<150> 60/294,823

<151> 2001-05-31

<150> 60/304,868

<151> 2001-07-12

<160> 127

<170> PatentIn Ver. 2.1

<210> 1

<211> 2838

<212> DNA

<213> Homo sapiens

<400> 1

```
aactttatga agctatggga cttgacaaaa agtgatattt gagaagaaag tacgcagtgg 60
ttgggtgtttt ctttttttta ataaaggaat tgaattactt tgaacacctc ttccagctgt 120
gcattacaga taacgtcagg aagagtctct gctttacaga atcggatttc atcacatgac 180
aacatgaagc tgtggattca tctcttttat tcatctctcc ttgcctgtat atctttacac 240
tcccaaactc cagtgtcttc atccagaggc tcttgtgatt ctctttgcaa ttgtgaggaa 300
aaagatggca caatgctaataa aaattgtgaa gcaaaaggta tcaagatggg atctgaaata 360
agtgtgccac catcacgacc tttccaacta agcttattaa ataacggctt gacgatgctt 420
cacacaaatg acttttctgg gcttaccaat gctatttcaa tacaccttgg atttaacaat 480
attgcagata ttgagatagg tgcatttaat ggccttggcc tcctgaaaca acttcataatc 540
aatcacaatt ctttagaaat tcttaaagag gatactttcc atggactgga aaacctggaa 600
ttcctgcaag cagataacaa ttttatcaca gtgattgaac caagtgcctt tagcaagctc 660
aacagactca aagtgttaat tttaaatgac aatgctattg agagtcttcc tccaaacatc 720
ttccgatttg ttcccttaac ccatctagat cttcgtggaa atcaattaca aacattgcct 780
tatgttggtt ttctcgaaca cattggccga atattggatc ttcagttgga ggacaacaaa 840
tgggcctgca attgtgactt attgcagtta aaaacttggg tggagaacat gcctccacag 900
tctataattg gtgatgttgt ctgcaacagc cctccatttt ttaaaggaag tatactcagt 960
agactaaaga aggaatctat ttgccctact ccaccagtgt atgaagaaca tgaggatcct 1020
tcaggatcat tacatttggc agcaacatct tcaataaatg atagtgcgat gtcaactaag 1080
accacgtcca ttctaataat acccaccaaa gcaccagggt tgatacctta tattacaaag 1140
ccatccactc aacttccagg accttactgc cctattcctt gtaactgcaa agtcctatcc 1200
ccatcaggac ttctaataca ttgtcaggag cgcaacattg aaagcttatc agatctgaga 1260
cctcctccgc aaaatcctag aaagctcatt ctagcgggaa atattattca cagtttaatg 1320
aagtctgac tagtggaata tttcactttg gaaatgcctt acttgggaaa caatcgtatt 1380
gaagttcttg aagaaggatc gtttatgaac ctaacgagat taaaaaaact ctatctaaat 1440
ggtaaccacc tgaccaaatt aagtaaaggc atgttccttg gtctccataa tcttgaatac 1500
ttatatcttg aatacaatgc cattaaggaa atactgccag gaacctttaa tccaatgcct 1560
aaacttaaag tctgtatatt aaataacaac ctctccaaag ttttaccacc acatattttt 1620
tcaggggttc ctctaactaa ggtaaatctt aaaacaaacc agtttaccga tctacctgta 1680
agtaatatct tggatgatct tgatttacta acccagattg accttgagga taacccttgg 1740
gactgtcctt gtgacctggg tggactgcag caatggatac aaaagttaag caagaacaca 1800
gtgacagatg acatcctctg cacttcccc gggtcatctg acaaaaagga attgaaagcc 1860
ctaaatagtg aaattctctg tccagggtta gtaaataacc catccatgcc aacacagact 1920
agttacctta tggtcaccac tcttgcacaa acaacaaata cggctgatac tattttacga 1980
tctcttacgg acgtgtgtgc actgtctgtt ctaatatggg gacttctgat tatgttcac 2040
actattgttt tctgtgtgtg agggatagtg gttctgttcc ttcaccgcag gagaagatac 2100
aaaaagaaac aagtagatga gcaaatgaga gacaacagtc ctgtgcatct tcagtacagc 2160
```

```

atgtatggcc ataaaaccac tcatcacact actgaaagac cctctgcctc actctatgaa 2220
cagcacatgg tgagcccat ggttcatgtc tatagaagtc catcctttgg tccaaagcat 2280
ctggaagagg aagaagagag gaatgagaaa gaaggaagtg atgcaaaaaca tctccaaaga 2340
agtcttttgg aacaggaaaa tcattcacca ctcacagggt caaatatgaa atacaaaacc 2400
acgaaccaat caacagaatt tttatccttc caagatgcc a gctcattgta cagaaacatt 2460
ttagaaaaag aaagggaact tcagcaactg ggaatcacag aatacctaag gaaaaacatt 2520
gctcagctcc agcctgatat ggaggcacat tctcctggag cccacgaaga gctgaagtta 2580
atggaaacat taatgtactc acgtccaagg aaggtattag tggaacagac aaaaaatgag 2640
tattttgaac ttaaagctaa tttacatgct gaacctgact atttagaagt cctggagcag 2700
caaacataga tggagagttt gagggctttc gcagaaatgc tgtgattctg ttttaagtcc 2760
ataccttgta aattagtgcc ttacgtgagt gtgtcatcca tcagaaccta agcacagcag 2820
taaactatgg agaaaaaa
2838

```

<210> 2
 <211> 841
 <212> PRT
 <213> Homo sapiens

```

<400> 2
Met Lys Leu Trp Ile His Leu Phe Tyr Ser Ser Leu Leu Ala Cys Ile
  1             5             10             15

Ser Leu His Ser Gln Thr Pro Val Leu Ser Ser Arg Gly Ser Cys Asp
      20             25             30

Ser Leu Cys Asn Cys Glu Glu Lys Asp Gly Thr Met Leu Ile Asn Cys
      35             40             45

Glu Ala Lys Gly Ile Lys Met Val Ser Glu Ile Ser Val Pro Pro Ser
      50             55             60

Arg Pro Phe Gln Leu Ser Leu Leu Asn Asn Gly Leu Thr Met Leu His
      65             70             75             80

Thr Asn Asp Phe Ser Gly Leu Thr Asn Ala Ile Ser Ile His Leu Gly
      85             90             95

Phe Asn Asn Ile Ala Asp Ile Glu Ile Gly Ala Phe Asn Gly Leu Gly
      100            105            110

Leu Leu Lys Gln Leu His Ile Asn His Asn Ser Leu Glu Ile Leu Lys
      115            120            125

Glu Asp Thr Phe His Gly Leu Glu Asn Leu Glu Phe Leu Gln Ala Asp
      130            135            140

Asn Asn Phe Ile Thr Val Ile Glu Pro Ser Ala Phe Ser Lys Leu Asn
      145            150            155            160

Arg Leu Lys Val Leu Ile Leu Asn Asp Asn Ala Ile Glu Ser Leu Pro
      165            170            175

Pro Asn Ile Phe Arg Phe Val Pro Leu Thr His Leu Asp Leu Arg Gly
      180            185            190

Asn Gln Leu Gln Thr Leu Pro Tyr Val Gly Phe Leu Glu His Ile Gly

```

195	200	205
Arg Ile Leu Asp Leu Gln Leu Glu Asp Asn Lys Trp Ala Cys Asn Cys 210 215 220		
Asp Leu Leu Gln Leu Lys Thr Trp Leu Glu Asn Met Pro Pro Gln Ser 225 230 235 240		
Ile Ile Gly Asp Val Val Cys Asn Ser Pro Pro Phe Phe Lys Gly Ser 245 250 255		
Ile Leu Ser Arg Leu Lys Lys Glu Ser Ile Cys Pro Thr Pro Pro Val 260 265 270		
Tyr Glu Glu His Glu Asp Pro Ser Gly Ser Leu His Leu Ala Ala Thr 275 280 285		
Ser Ser Ile Asn Asp Ser Arg Met Ser Thr Lys Thr Thr Ser Ile Leu 290 295 300		
Lys Leu Pro Thr Lys Ala Pro Gly Leu Ile Pro Tyr Ile Thr Lys Pro 305 310 315 320		
Ser Thr Gln Leu Pro Gly Pro Tyr Cys Pro Ile Pro Cys Asn Cys Lys 325 330 335		
Val Leu Ser Pro Ser Gly Leu Leu Ile His Cys Gln Glu Arg Asn Ile 340 345 350		
Glu Ser Leu Ser Asp Leu Arg Pro Pro Pro Gln Asn Pro Arg Lys Leu 355 360 365		
Ile Leu Ala Gly Asn Ile Ile His Ser Leu Met Lys Ser Asp Leu Val 370 375 380		
Glu Tyr Phe Thr Leu Glu Met Leu His Leu Gly Asn Asn Arg Ile Glu 385 390 395 400		
Val Leu Glu Glu Gly Ser Phe Met Asn Leu Thr Arg Leu Gln Lys Leu 405 410 415		
Tyr Leu Asn Gly Asn His Leu Thr Lys Leu Ser Lys Gly Met Phe Leu 420 425 430		
Gly Leu His Asn Leu Glu Tyr Leu Tyr Leu Glu Tyr Asn Ala Ile Lys 435 440 445		
Glu Ile Leu Pro Gly Thr Phe Asn Pro Met Pro Lys Leu Lys Val Leu 450 455 460		
Tyr Leu Asn Asn Asn Leu Leu Gln Val Leu Pro Pro His Ile Phe Ser 465 470 475 480		
Gly Val Pro Leu Thr Lys Val Asn Leu Lys Thr Asn Gln Phe Thr His 485 490 495		
Leu Pro Val Ser Asn Ile Leu Asp Asp Leu Asp Leu Leu Thr Gln Ile		

500	505	510
Asp Leu Glu Asp Asn Pro Trp	Asp Cys Ser Cys Asp	Leu Val Gly Leu
515	520	525
Gln Gln Trp Ile Gln Lys Leu Ser Lys Asn Thr Val Thr Asp Asp Ile		
530	535	540
Leu Cys Thr Ser Pro Gly His Leu Asp Lys Lys Glu Leu Lys Ala Leu		
545	550	555 560
Asn Ser Glu Ile Leu Cys Pro Gly Leu Val Asn Asn Pro Ser Met Pro		
565	570	575
Thr Gln Thr Ser Tyr Leu Met Val Thr Thr Pro Ala Thr Thr Thr Asn		
580	585	590
Thr Ala Asp Thr Ile Leu Arg Ser Leu Thr Asp Ala Val Pro Leu Ser		
595	600	605
Val Leu Ile Leu Gly Leu Leu Ile Met Phe Ile Thr Ile Val Phe Cys		
610	615	620
Ala Ala Gly Ile Val Val Leu Val Leu His Arg Arg Arg Arg Tyr Lys		
625	630	635 640
Lys Lys Gln Val Asp Glu Gln Met Arg Asp Asn Ser Pro Val His Leu		
645	650	655
Gln Tyr Ser Met Tyr Gly His Lys Thr Thr His His Thr Thr Glu Arg		
660	665	670
Pro Ser Ala Ser Leu Tyr Glu Gln His Met Val Ser Pro Met Val His		
675	680	685
Val Tyr Arg Ser Pro Ser Phe Gly Pro Lys His Leu Glu Glu Glu Glu		
690	695	700
Glu Arg Asn Glu Lys Glu Gly Ser Asp Ala Lys His Leu Gln Arg Ser		
705	710	715 720
Leu Leu Glu Gln Glu Asn His Ser Pro Leu Thr Gly Ser Asn Met Lys		
725	730	735
Tyr Lys Thr Thr Asn Gln Ser Thr Glu Phe Leu Ser Phe Gln Asp Ala		
740	745	750
Ser Ser Leu Tyr Arg Asn Ile Leu Glu Lys Glu Arg Glu Leu Gln Gln		
755	760	765
Leu Gly Ile Thr Glu Tyr Leu Arg Lys Asn Ile Ala Gln Leu Gln Pro		
770	775	780
Asp Met Glu Ala His Tyr Pro Gly Ala His Glu Glu Leu Lys Leu Met		
785	790	795 800
Glu Thr Leu Met Tyr Ser Arg Pro Arg Lys Val Leu Val Glu Gln Thr		

805

810

815

Lys Asn Glu Tyr Phe Glu Leu Lys Ala Asn Leu His Ala Glu Pro Asp
 820 825 830

Tyr Leu Glu Val Leu Glu Gln Gln Thr
 835 840

<210> 3

<211> 2526

<212> DNA

<213> Homo sapiens

<400> 3

```

atgaagctgt ggattcatct cttttattca tctctccttg cctgtatatac tttacactcc 60
caaactccag tgctctcatc cagaggctct tgtgattctc tttgcaattg tgaggaaaaa 120
gatggcaca tgctaataaa ttgtgaagca aaaggatatc agatgggtatc tgaaataagt 180
gtgccaccat caccaccttt ccaactaagc ttattaaata acggcttgac gatgcttcac 240
acaaatgact tttctgggct taccaatgct atttcaatac accttggatt taacaatatt 300
gcagatattg agataggtgc atttaatggc cttggcctcc tgaaacaact tcatatcaat 360
cacaattctt tagaaattct taaagaggat actttccatg gactggaaaa cctggaattc 420
ctgcaagcag ataacaattt tatcacagt attgaaccaa gtgccttttag caagctcaac 480
agactcaaag tgtaattttt aaatgacaat gctattgaga gtcttcctcc aaacatcttc 540
cgatttggtc ctttaaccca tctagatctt cgtggaaatc aattacaaac attgccttat 600
gttggttttc tcgaacacat tggccgaata ttggatcttc agttggagga caacaaatgg 660
gcctgcaatt gtgacttatt gcagttaaaa acttgggttg agaacatgcc tccacagtct 720
ataattggtg atgttgtctg caacagccct ccatttttta aaggaagtat actcagtaga 780
ctaaagaagg aatctatttg ccctactcca ccagtgtatg aagaacatga ggatccttca 840
ggatcattac atctggcagc aacatcttca ataaatgata gtcgcatgtc aactaagacc 900
acgtccattc taaaactacc caccaaagca ccaggtttga taccttatat taaaaagcca 960
tccactcaac ttccaggacc ttactgccct attccttgta actgcaaagt cctatcccca 1020
tcaggacttc taatacattg tcaggagcgc aacattgaaa gcttatcaga tctgagacct 1080
cctccgcaaa atcctagaaa gctcattcta gcgggaaata ttattcacag tttaatgaag 1140
tctgatctag tggaatattt cactttggaa atgcttactc tgggaaacaa tcgtattgaa 1200
gttcttgaag aaggatcggt tatgaacctc acgagattac aaaaactcta tctaaatggt 1260
aaccacctga ccaaattaag taaaggcatg ttccttggtc tccataatct tgaatactta 1320
tatcttgaat acaatgccat taaggaaata ctgccaggaa cctttaatcc aatgcctaaa 1380
cttaaagtc tgatttaaaa taacaacctc ctccaagttt taccaccaca tattttttca 1440
ggggttcctc taactaaggt aaatcttaaa acaaaccagt ttaccatctt acctgtaagt 1500
aatatttttg atgatcttga tttgctaacc cagattgacc ttgaggataa cccctgggac 1560
tgctcctgtg acctggttgg actgcagcaa tggatacaaa agttaagcaa gaacacagt 1620
acagatgaca tcctctgcac ttccccggg catctcgaca aaaaggaatt gaaagcccta 1680
aatagtgaat ttctctgtcc aggttttagta aataaccat ccattgccaac acagactagt 1740
taccttatgg tcaccactcc tgcaacaaca acaatacgg ctgatactat tttacgatct 1800
cttacggacg ctgtgccact gtctgttcta atattgggac ttctgattat gttcatcact 1860
attgttttct gtgctgcagg gatagtgggt cttgttcttc accgcaggag aagatacaaa 1920
aagaaacaag tagatgagca aatgagagac aacagtcctg tgcattctca gtacagcatg 1980
tatggccata aaaccactca tcacactact gaaagacct ctgcctcact ctatgaacag 2040
cacatggtga gccccatggt tcatgtctat agaagtccat ccttttggtc aaagcatctg 2100
gaagaggaag aagagaggaa tgagaaagaa ggaagtgatg caaaacatct ccaaagaagt 2160
cttttggaac aggaaaatca ttcaccactc acaggggtcaa atatgaaata caaaaccacg 2220
aaccaatcaa cagaattttt atccttccaa gatgccagct cattgtacag aaacattttta 2280
gaaaaagaaa gggaacttca gcaactggga atcacagaat acctaaggaa aaacattgct 2340
cagctccagc ctgatatgga ggcacattat cctggagccc acgaagagct gaagttaagt 2400
gaaacattaa tgtactcacg tccaaggaag gtattagtgg aacagacaaa aaatgagtat 2460
tttgaactta aagctaattt acatgctgaa cctgactatt tagaagtcct ggagcagcaa 2520

```

<210> 4

<211> 2531

<212> DNA

<213> Homo sapiens

<400> 4

```

ggattctctc ttttattcat ctctccttgc ctgtatatct ttacactccc aaactccagt 60
gctctcatcc agaggctctt gtgattctct ttgcaattgt gaggaaaaag atggcacaat 120
gctaataaat tgtgaagcaa aagggtatcaa gatggtatct gaaataagtg tgctaccatc 180
acgacctttc caactaagct tattaaataa cggcttgacg atgcttcaca caaatgactt 240
ttctgggctt accaatgcta tttcaatata ccttggattt aacaatattg cagatattga 300
gatagggtgca tttaatggcc ttggcctcct gaaacaactt catatcaatc acaattcctt 360
agaaattctt aaagaggata ctttccatgg actggaaaac ctggaattcc tgcaagcaga 420
taacaatttt atcacagtga ttgaaccaag tgcctttagc aagctcaaca gactcaaagt 480
gttaatttta aatgacaatg ctattgagag tcttcctcca aacatcttcc gatttgttcc 540
tttaacccat ctagatcttc gtggaaatca attacaaaca ttgccttatg ttggttttct 600
cgaacacatt ggccgaatat tggatcttca gttggaggac aacaaatggg cctgcaattg 660
tgacttattg cagttaaaaa cttgggttga gaacatgcct ccacagtcta taattggtga 720
tggtgtctgc aacagccctc cattttttta aggaagtata ctacgtagac taaagaagga 780
atctattttg cctactccac cagtgtatga agaacatgag gatccttcag gatcattaca 840
tctggcagca acatcttcaa taaatgatag tcgcagtgtc actaagacca cgtccattct 900
aaaactaccc accaaagcac caggtttgat accttatatt acaaagccat ccactcaact 960
tccaggacct tactgcccta ttccttgtaa ctgcaaagtc ctatcccat caggacttct 1020
aatacattgt caggagcgca acattgaaag cttatcagat ctgagacctc ctccgcaaaa 1080
tcctagaaag ctcatcttag cgggaaatat tattcacagt ttaatgaagt ctgatctagt 1140
ggaatatttc actttgaaa tgcttcactt gggaaacaat cgtattgaag ttcttgaaga 1200
aggatcggtt atgaacctaa cgagattaca aaaactctat ctaaatggta accacctgac 1260
caaattaagt aaaggcatgt tccttgggtc ccataatctt gaatacttat atcttgaata 1320
caatgccatt aaggaaatac tgccaggaac ctttaatcca atgcctaaac ttaaagtcct 1380
gtattttaat aacaacctcc tccaagtttt accaccacat attttttcag gggttcctct 1440
aactaaggta aatcttaaaa caaaccagtt taccatctc cctgtaagta atattttgga 1500
tgatcttgat ttgctaacc agattgacct tgaggataac ccctgggact gtcctgtga 1560
cctgggttga ctgcagcaat ggatacaaaa gttaagcaag aacacagtga cagatgacat 1620
cctctgcact tccccgggc atctcgacaa aaaggaattg aaagccctaa atagtgaat 1680
tctctgtcca ggtttagtaa ataaccatc catgccaaca cagactagtt acctatggt 1740
caccactcct gcaacaacaa caaatacggc tgatactatt ttacgatctc ttacggacgc 1800
tgtgccactg tctgttctaa tattgggact tctgattatg ttcactacta ttgttttctg 1860
tgctgcaggg atagtgggtc ttgttcttca ccgcaggaga agatacaaaa agaaacaagt 1920
agatgagcaa atgagagaca acagtcctgt gcatcttcag tacagcatgt atggccataa 1980
aaccactcat cacactactg aaagacctc tgccctactc tatgaacagc acatgggtgag 2040
ccccatggtt catgtctata gaagtccatc ctttgggtcca aagcatctgg aagaggaaga 2100
agagaggaat gagaaagaag gaagtgatgc aaaacatctc caaagaagtc ttttgaaca 2160
ggaaaatcat tcaccactca cagggtcaaa tatgaaatac aaaaccacga accaatcaac 2220
agaattttta tccttccaag atgccagctc attgtacaga aacatttttag aaaaagaaag 2280
ggaacttcag caactgggaa tcacagaata cctaaggaaa aacattgctc agctccagcc 2340
tgatatggag gcacattatc ctggagccca cgaagagctg aagttaatgg aaacattaat 2400
gtactcacgt ccaaggaagg tattagtggg acagacaaaa aatgagtatt ttgaacttaa 2460
agctaattta catgctgaac ctgactattt agaagtcctg gagcagcaaa cataagggcg 2520
aattctgctg t

```

2531

<210> 5

<211> 837

<212> PRT

<213> Homo sapiens

<400> 5

Asp Ser Leu Phe Tyr Ser Ser Leu Leu Ala Cys Ile Ser Leu His Ser
1 5 10 15
Gln Thr Pro Val Leu Ser Ser Arg Gly Ser Cys Asp Ser Leu Cys Asn
20 25 30
Cys Glu Glu Lys Asp Gly Thr Met Leu Ile Asn Cys Glu Ala Lys Gly
35 40 45
Ile Lys Met Val Ser Glu Ile Ser Val Leu Pro Ser Arg Pro Phe Gln
50 55 60
Leu Ser Leu Leu Asn Asn Gly Leu Thr Met Leu His Thr Asn Asp Phe
65 70 75 80
Ser Gly Leu Thr Asn Ala Ile Ser Ile His Leu Gly Phe Asn Asn Ile
85 90 95
Ala Asp Ile Glu Ile Gly Ala Phe Asn Gly Leu Gly Leu Leu Lys Gln
100 105 110
Leu His Ile Asn His Asn Ser Leu Glu Ile Leu Lys Glu Asp Thr Phe
115 120 125
His Gly Leu Glu Asn Leu Glu Phe Leu Gln Ala Asp Asn Asn Phe Ile
130 135 140
Thr Val Ile Glu Pro Ser Ala Phe Ser Lys Leu Asn Arg Leu Lys Val
145 150 155 160
Leu Ile Leu Asn Asp Asn Ala Ile Glu Ser Leu Pro Pro Asn Ile Phe
165 170 175
Arg Phe Val Pro Leu Thr His Leu Asp Leu Arg Gly Asn Gln Leu Gln
180 185 190
Thr Leu Pro Tyr Val Gly Phe Leu Glu His Ile Gly Arg Ile Leu Asp
195 200 205
Leu Gln Leu Glu Asp Asn Lys Trp Ala Cys Asn Cys Asp Leu Leu Gln
210 215 220
Leu Lys Thr Trp Leu Glu Asn Met Pro Pro Gln Ser Ile Ile Gly Asp
225 230 235 240
Val Val Cys Asn Ser Pro Pro Phe Phe Lys Gly Ser Ile Leu Ser Arg
245 250 255
Leu Lys Lys Glu Ser Ile Cys Pro Thr Pro Pro Val Tyr Glu Glu His
260 265 270
Glu Asp Pro Ser Gly Ser Leu His Leu Ala Ala Thr Ser Ser Ile Asn
275 280 285

Asp Ser Arg Met Ser Thr Lys Thr Thr Ser Ile Leu Lys Leu Pro Thr
 290 295 300
 Lys Ala Pro Gly Leu Ile Pro Tyr Ile Thr Lys Pro Ser Thr Gln Leu
 305 310 315 320
 Pro Gly Pro Tyr Cys Pro Ile Pro Cys Asn Cys Lys Val Leu Ser Pro
 325 330 335
 Ser Gly Leu Leu Ile His Cys Gln Glu Arg Asn Ile Glu Ser Leu Ser
 340 345 350
 Asp Leu Arg Pro Pro Pro Gln Asn Pro Arg Lys Leu Ile Leu Ala Gly
 355 360 365
 Asn Ile Ile His Ser Leu Met Lys Ser Asp Leu Val Glu Tyr Phe Thr
 370 375 380
 Leu Glu Met Leu His Leu Gly Asn Asn Arg Ile Glu Val Leu Glu Glu
 385 390 395 400
 Gly Ser Phe Met Asn Leu Thr Arg Leu Gln Lys Leu Tyr Leu Asn Gly
 405 410 415
 Asn His Leu Thr Lys Leu Ser Lys Gly Met Phe Leu Gly Leu His Asn
 420 425 430
 Leu Glu Tyr Leu Tyr Leu Glu Tyr Asn Ala Ile Lys Glu Ile Leu Pro
 435 440 445
 Gly Thr Phe Asn Pro Met Pro Lys Leu Lys Val Leu Tyr Leu Asn Asn
 450 455 460
 Asn Leu Leu Gln Val Leu Pro Pro His Ile Phe Ser Gly Val Pro Leu
 465 470 475 480
 Thr Lys Val Asn Leu Lys Thr Asn Gln Phe Thr His Leu Pro Val Ser
 485 490 495
 Asn Ile Leu Asp Asp Leu Asp Leu Leu Thr Gln Ile Asp Leu Glu Asp
 500 505 510
 Asn Pro Trp Asp Cys Ser Cys Asp Leu Val Gly Leu Gln Gln Trp Ile
 515 520 525
 Gln Lys Leu Ser Lys Asn Thr Val Thr Asp Asp Ile Leu Cys Thr Ser
 530 535 540
 Pro Gly His Leu Asp Lys Lys Glu Leu Lys Ala Leu Asn Ser Glu Ile
 545 550 555 560
 Leu Cys Pro Gly Leu Val Asn Asn Pro Ser Met Pro Thr Gln Thr Ser
 565 570 575
 Tyr Leu Met Val Thr Thr Pro Ala Thr Thr Thr Asn Thr Ala Asp Thr
 580 585 590

Ile Leu Arg Ser Leu Thr Asp Ala Val Pro Leu Ser Val Leu Ile Leu
 595 600 605
 Gly Leu Leu Ile Met Phe Ile Thr Ile Val Phe Cys Ala Ala Gly Ile
 610 615 620
 Val Val Leu Val Leu His Arg Arg Arg Arg Tyr Lys Lys Lys Gln Val
 625 630 635 640
 Asp Glu Gln Met Arg Asp Asn Ser Pro Val His Leu Gln Tyr Ser Met
 645 650 655
 Tyr Gly His Lys Thr Thr His His Thr Thr Glu Arg Pro Ser Ala Ser
 660 665 670
 Leu Tyr Glu Gln His Met Val Ser Pro Met Val His Val Tyr Arg Ser
 675 680 685
 Pro Ser Phe Gly Pro Lys His Leu Glu Glu Glu Glu Glu Arg Asn Glu
 690 695 700
 Lys Glu Gly Ser Asp Ala Lys His Leu Gln Arg Ser Leu Leu Glu Gln
 705 710 715 720
 Glu Asn His Ser Pro Leu Thr Gly Ser Asn Met Lys Tyr Lys Thr Thr
 725 730 735
 Asn Gln Ser Thr Glu Phe Leu Ser Phe Gln Asp Ala Ser Ser Leu Tyr
 740 745 750
 Arg Asn Ile Leu Glu Lys Glu Arg Glu Leu Gln Gln Leu Gly Ile Thr
 755 760 765
 Glu Tyr Leu Arg Lys Asn Ile Ala Gln Leu Gln Pro Asp Met Glu Ala
 770 775 780
 His Tyr Pro Gly Ala His Glu Glu Leu Lys Leu Met Glu Thr Leu Met
 785 790 795 800
 Tyr Ser Arg Pro Arg Lys Val Leu Val Glu Gln Thr Lys Asn Glu Tyr
 805 810 815
 Phe Glu Leu Lys Ala Asn Leu His Ala Glu Pro Asp Tyr Leu Glu Val
 820 825 830
 Leu Glu Gln Gln Thr
 835

<210> 6

<211> 3609

<212> DNA

<213> Homo sapiens

<400> 6

cagtggatgc agaaggcaga cagcagcacc gagacgatga aggagaagag gacagcggct 60
 gcgatcaccg tgcggcacag gaccggctcc tgcttctcgg gccgctgtgt caactccacc 120

tgcctctgcg	acccgggctg	ggtgggggac	cagtgccagc	actgccaggg	caggttcagg	180
ttaacagaac	cttctggata	tttaacagat	ggcccaatta	actataaata	taaaactaaa	240
tgtacttgge	tcattgaagg	cccaaagtca	gtgttaagat	taagattcaa	tcattttgct	300
acagaatgta	gctgggatca	tatgtatggt	tatgatggag	attcaatata	tgacacctta	360
atagcttctt	ttagtgggtt	gatagtcctt	gaaataaggg	gcaatgaaac	tgtgcctgaa	420
gttggtacta	catctggcta	tgactgttta	cattttttta	gtgatgctgc	gtataatcta	480
actggtttca	acatttttcta	ttcgatcaat	tcttgtccta	acaattgctc	tggcatggg	540
aagtgtacaa	ctagtgtctc	tgttccaagt	caagtatatt	gtgaatgtga	taaatactgg	600
aagggtgaag	cttgtgatat	tccttactgt	aaagccaatt	gcggcagtcc	agatcacggg	660
tactgtgacc	tgactggaga	aaaattatgt	gtctgcaatg	atagttggca	aggtataggt	720
cctgattggt	ctttgaatgt	tccttctact	gagtccttact	ggattctgcc	aaacgttaaa	780
cccttcagtc	cttctgtagg	tcgggcttca	cataaagcag	ttttacacgg	gaaatttatg	840
tgggtgattg	gtggatatac	ttttaactac	agttcttttc	aaatggctct	aagttacaat	900
ttagaaagca	gtatatggaa	tgtaggaact	ccatcaaggg	gacctctcca	gagatatgga	960
cactctcttg	ctttatatca	ggaaaacatc	tttatgtatg	gaggcagaat	tgaacaaaat	1020
gatggcaatg	tcacagatga	attatgggtt	tttaacatac	atagtcagtc	atggagtaca	1080
aaaactccta	ctgttcttgg	acatggtcag	cagtatgctg	tggagggaca	ttcagcacat	1140
attatggagt	tggatagtag	agatgttgct	atgatcataa	tatttggata	ttctgcaata	1200
tattggttata	caagcagcat	acaggaatac	catatctgtt	caaacacttg	gcttgttcca	1260
gaaactaaag	gagctattgt	acaaggtgga	tatggccata	ctagtgtgta	tgatgaaata	1320
acaaagtcca	tttatgttca	tggaggggat	aaagcattgc	cagggaacaa	atatggattg	1380
gttgatgata	tttataaata	tgaagttaac	actaagactt	ggactatttt	gaaagaaagt	1440
gggtttgcca	gataccttca	ttcagctggt	cttatcaatg	gagctatgct	tatttttggg	1500
ggaaataccc	ataatgacac	ttccttgagt	aacgggtgca	aatgtttttc	tgccgattttc	1560
ctggcatatg	acatatgccc	aggctggagt	gcagtggcac	gatctcagct	cactgccacc	1620
tcacactccc	acgttcaagc	gattctcaat	aggctccatg	atatatttgg	gggattttct	1680
agtgtactcc	ttaatgatat	ccttgatac	aagcctccaa	attgcaaggc	tttcagagat	1740
gaagaacttt	gtaaaaatgc	tggtccaggg	ataaaatgtg	tttgaataa	aaatcactgt	1800
gaatcttggg	aatctgggaa	tactaataat	attcttagag	caaagtgcct	ttctaaaaga	1860
aatctctgca	gtgacagatg	ttacagatat	gcagattgtg	ccagctgtac	tgccaatata	1920
aatgggtgcc	aatgggtgtg	tgacaagaaa	tgcatttcgg	caaatagtaa	ctgcagtatg	1980
gttagtattt	ttgggtatat	aaccttgcct	tcacagttcc	cattctatta	ttgctacaga	2040
tatgcagatt	gtgccagctg	tactgccaat	acaaatgggt	gccaatggtg	tgatgacaag	2100
aaatgcattg	ctttaccagc	tcattcttgt	ggagaaggat	ggagtcatat	tggggatgct	2160
tgtcttagag	tcaattccag	tagagaaaac	tatgacaatg	caaaacttta	ttgctataat	2220
cttagtgga	atcttgcttc	attaacaacc	tcaaaagaag	tagaatttgt	tctggatgaa	2280
atacagaagt	atacacaaca	gaaagtatca	ccttgggtag	gcttgcgcaa	gatcaatata	2340
tcctattggg	gatgggaaga	catgtctcct	tttacaacaa	caacactaca	gtggcttcct	2400
ggcgaaccca	atgattctgg	gttttgtgca	tatctggaaa	gggctgcagt	ggcaggctta	2460
aaagctaata	cttgtagatc	tatggcaaat	ggccttgtct	gtgaaaaacc	tgtaaatcaa	2520
aatgcgaggc	cgtgcaaaaa	gccatgctct	ctgaggacat	catgttccaa	ctgtacaagc	2580
aatggcatgg	agtgtatgtg	gtgcagcagt	acgaaacgat	gtgttgactc	taatgcctat	2640
atcatctctt	ttccatatgg	acaatgtcta	gagtggcaaa	ctgccacctg	ctcccgctgt	2700
caaaattggt	ctggattgag	aacctgtgga	cagtgttttg	aacagcctga	atgtggctgg	2760
tgcaatgatc	ctagtaatac	aggaagagga	cattgcattg	aaggttcttc	acggggacca	2820
atgaagctta	ttggaatgca	ccacagtggg	atggttcttg	acaccaatct	ttgccccaaa	2880
gaaaagaact	atgagtgggtc	ctttatccag	tgtccagctt	gccagtgtaa	tggacatagc	2940
acttgcata	ataataatgt	gtgcgaacag	tgtaaaaatc	tcaccacagg	aaagcagtgt	3000
caagattgta	tgccagggtta	ttatggagat	ccaaccaatg	gtggacagtg	cacagcttgt	3060
acatgcagtg	gccatgcaaa	tatctgtcat	ctgcacacag	gaaaatgttt	ctgcacaact	3120
aaaggaataa	aagggtgacca	atgccaatgt	tgtgactctg	aaaatcgcta	tgttggtaat	3180
ccacttagag	gaacatgtta	ttgtaagtat	agccttttga	ttgattatca	atttaccttc	3240
agcttattac	aggaagatga	tcgccaccat	actgccataa	acttttatagc	aaaccagaa	3300
caggtgagga	aaaatctgga	tatatcaatt	aatgcatcaa	acaactttta	tctcaacatt	3360
acgtggtctg	tcggttcagc	tggaaacaata	tctggggaag	agacttctat	agtttccaag	3420
aataatataa	aggaatacag	agatagtttt	tcctatgaaa	aattttaactt	tagaagcaat	3480
cctaacatta	cattctatgt	gtacgtcagc	aacttttctt	ggcctattta	aatacaggta	3540

agtgtaaga gtatttactt ctaatgacca taatatcatt aagaaaagaa tgggtgctttt 3600
gtccaaagt 3609

<210> 7
<211> 1185
<212> PRT
<213> Homo sapiens

<400> 7

Met	Gln	Lys	Ala	Asp	Ser	Ser	Thr	Glu	Thr	Met	Lys	Glu	Lys	Arg	Thr	1	5	10	15
Ala	Ala	Ala	Ile	Thr	Val	Arg	His	Arg	Thr	Gly	Ser	Cys	Phe	Ser	Gly	20	25	30	
Arg	Cys	Val	Asn	Ser	Thr	Cys	Leu	Cys	Asp	Pro	Gly	Trp	Val	Gly	Asp	35	40	45	
Gln	Cys	Gln	His	Cys	Gln	Gly	Arg	Phe	Arg	Leu	Thr	Glu	Pro	Ser	Gly	50	55	60	
Tyr	Leu	Thr	Asp	Gly	Pro	Ile	Asn	Tyr	Lys	Tyr	Lys	Thr	Lys	Cys	Thr	65	70	75	80
Trp	Leu	Ile	Glu	Gly	Pro	Asn	Ala	Val	Leu	Arg	Leu	Arg	Phe	Asn	His	85	90	95	
Phe	Ala	Thr	Glu	Cys	Ser	Trp	Asp	His	Met	Tyr	Val	Tyr	Asp	Gly	Asp	100	105	110	
Ser	Ile	Tyr	Ala	Pro	Leu	Ile	Ala	Ser	Phe	Ser	Gly	Leu	Ile	Val	Pro	115	120	125	
Glu	Ile	Arg	Gly	Asn	Glu	Thr	Val	Pro	Glu	Val	Val	Thr	Thr	Ser	Gly	130	135	140	
Tyr	Ala	Leu	Leu	His	Phe	Phe	Ser	Asp	Ala	Ala	Tyr	Asn	Leu	Thr	Gly	145	150	155	160
Phe	Asn	Ile	Phe	Tyr	Ser	Ile	Asn	Ser	Cys	Pro	Asn	Asn	Cys	Ser	Gly	165	170	175	
His	Gly	Lys	Cys	Thr	Thr	Ser	Val	Ser	Val	Pro	Ser	Gln	Val	Tyr	Cys	180	185	190	
Glu	Cys	Asp	Lys	Tyr	Trp	Lys	Gly	Glu	Ala	Cys	Asp	Ile	Pro	Tyr	Cys	195	200	205	
Lys	Ala	Asn	Cys	Gly	Ser	Pro	Asp	His	Gly	Tyr	Cys	Asp	Leu	Thr	Gly	210	215	220	
Glu	Lys	Leu	Cys	Val	Cys	Asn	Asp	Ser	Trp	Gln	Gly	Ile	Gly	Pro	Asp	225	230	235	240
Cys	Ser	Leu	Asn	Val	Pro	Ser	Thr	Glu	Ser	Tyr	Trp	Ile	Leu	Pro	Asn	245	250	255	

Val Lys Pro Phe Ser Pro Ser Val Gly Arg Ala Ser His Lys Ala Val
 260 265 270
 Leu His Gly Lys Phe Met Trp Val Ile Gly Gly Tyr Thr Phe Asn Tyr
 275 280 285
 Ser Ser Phe Gln Met Val Leu Ser Tyr Asn Leu Glu Ser Ser Ile Trp
 290 295 300
 Asn Val Gly Thr Pro Ser Arg Gly Pro Leu Gln Arg Tyr Gly His Ser
 305 310 315 320
 Leu Ala Leu Tyr Gln Glu Asn Ile Phe Met Tyr Gly Gly Arg Ile Glu
 325 330 335
 Thr Asn Asp Gly Asn Val Thr Asp Glu Leu Trp Val Phe Asn Ile His
 340 345 350
 Ser Gln Ser Trp Ser Thr Lys Thr Pro Thr Val Leu Gly His Gly Gln
 355 360 365
 Gln Tyr Ala Val Glu Gly His Ser Ala His Ile Met Glu Leu Asp Ser
 370 375 380
 Arg Asp Val Val Met Ile Ile Ile Phe Gly Tyr Ser Ala Ile Tyr Gly
 385 390 395 400
 Tyr Thr Ser Ser Ile Gln Glu Tyr His Ile Cys Ser Asn Thr Trp Leu
 405 410 415
 Val Pro Glu Thr Lys Gly Ala Ile Val Gln Gly Gly Tyr Gly His Thr
 420 425 430
 Ser Val Tyr Asp Glu Ile Thr Lys Ser Ile Tyr Val His Gly Gly Tyr
 435 440 445
 Lys Ala Leu Pro Gly Asn Lys Tyr Gly Leu Val Asp Asp Leu Tyr Lys
 450 455 460
 Tyr Glu Val Asn Thr Lys Thr Trp Thr Ile Leu Lys Glu Ser Gly Phe
 465 470 475 480
 Ala Arg Tyr Leu His Ser Ala Val Leu Ile Asn Gly Ala Met Leu Ile
 485 490 495
 Phe Gly Gly Asn Thr His Asn Asp Thr Ser Leu Ser Asn Gly Ala Lys
 500 505 510
 Cys Phe Ser Ala Asp Phe Leu Ala Tyr Asp Ile Cys Pro Gly Trp Ser
 515 520 525
 Ala Val Ala Arg Ser Gln Leu Thr Ala Thr Ser Thr Ser His Val Gln
 530 535 540
 Ala Ile Leu Asn Arg Ser Met Tyr Ile Phe Gly Gly Phe Ser Ser Val
 545 550 555 560

Leu Leu Asn Asp Ile Leu Val Tyr Lys Pro Pro Asn Cys Lys Ala Phe
 565 570 575
 Arg Asp Glu Glu Leu Cys Lys Asn Ala Gly Pro Gly Ile Lys Cys Val
 580 585 590
 Trp Asn Lys Asn His Cys Glu Ser Trp Glu Ser Gly Asn Thr Asn Asn
 595 600 605
 Ile Leu Arg Ala Lys Cys Phe Ser Lys Arg Asn Leu Cys Ser Asp Arg
 610 615 620
 Cys Tyr Arg Tyr Ala Asp Cys Ala Ser Cys Thr Ala Asn Thr Asn Gly
 625 630 635 640
 Cys Gln Trp Cys Asp Asp Lys Lys Cys Ile Ser Ala Asn Ser Asn Cys
 645 650 655
 Ser Met Val Ser Ile Phe Gly Tyr Ile Thr Leu Pro Ser Gln Phe Pro
 660 665 670
 Phe Tyr Tyr Cys Tyr Arg Tyr Ala Asp Cys Ala Ser Cys Thr Ala Asn
 675 680 685
 Thr Asn Gly Cys Gln Trp Cys Asp Asp Lys Lys Cys Ile Ala Leu Pro
 690 695 700
 Ala His Leu Cys Gly Glu Gly Trp Ser His Ile Gly Asp Ala Cys Leu
 705 710 715 720
 Arg Val Asn Ser Ser Arg Glu Asn Tyr Asp Asn Ala Lys Leu Tyr Cys
 725 730 735
 Tyr Asn Leu Ser Gly Asn Leu Ala Ser Leu Thr Thr Ser Lys Glu Val
 740 745 750
 Glu Phe Val Leu Asp Glu Ile Gln Lys Tyr Thr Gln Gln Lys Val Ser
 755 760 765
 Pro Trp Val Gly Leu Arg Lys Ile Asn Ile Ser Tyr Trp Gly Trp Glu
 770 775 780
 Asp Met Ser Pro Phe Thr Asn Thr Thr Leu Gln Trp Leu Pro Gly Glu
 785 790 795 800
 Pro Asn Asp Ser Gly Phe Cys Ala Tyr Leu Glu Arg Ala Ala Val Ala
 805 810 815
 Gly Leu Lys Ala Asn Pro Cys Thr Ser Met Ala Asn Gly Leu Val Cys
 820 825 830
 Glu Lys Pro Val Asn Gln Asn Ala Arg Pro Cys Lys Lys Pro Cys Ser
 835 840 845
 Leu Arg Thr Ser Cys Ser Asn Cys Thr Ser Asn Gly Met Glu Cys Met
 850 855 860

Trp Cys Ser Ser Thr Lys Arg Cys Val Asp Ser Asn Ala Tyr Ile Ile
 865 870 875 880
 Ser Phe Pro Tyr Gly Gln Cys Leu Glu Trp Gln Thr Ala Thr Cys Ser
 885 890 895
 Arg Ala Gln Asn Cys Ser Gly Leu Arg Thr Cys Gly Gln Cys Leu Glu
 900 905 910
 Gln Pro Glu Cys Gly Trp Cys Asn Asp Pro Ser Asn Thr Gly Arg Gly
 915 920 925
 His Cys Ile Glu Gly Ser Ser Arg Gly Pro Met Lys Leu Ile Gly Met
 930 935 940
 His His Ser Glu Met Val Leu Asp Thr Asn Leu Cys Pro Lys Glu Lys
 945 950 955 960
 Asn Tyr Glu Trp Ser Phe Ile Gln Cys Pro Ala Cys Gln Cys Asn Gly
 965 970 975
 His Ser Thr Cys Ile Asn Asn Asn Val Cys Glu Gln Cys Lys Asn Leu
 980 985 990
 Thr Thr Gly Lys Gln Cys Gln Asp Cys Met Pro Gly Tyr Tyr Gly Asp
 995 1000 1005
 Pro Thr Asn Gly Gly Gln Cys Thr Ala Cys Thr Cys Ser Gly His Ala
 1010 1015 1020
 Asn Ile Cys His Leu His Thr Gly Lys Cys Phe Cys Thr Thr Lys Gly
 1025 1030 1035 1040
 Ile Lys Gly Asp Gln Cys Gln Leu Cys Asp Ser Glu Asn Arg Tyr Val
 1045 1050 1055
 Gly Asn Pro Leu Arg Gly Thr Cys Tyr Cys Lys Tyr Ser Leu Leu Ile
 1060 1065 1070
 Asp Tyr Gln Phe Thr Phe Ser Leu Leu Gln Glu Asp Asp Arg His His
 1075 1080 1085
 Thr Ala Ile Asn Phe Ile Ala Asn Pro Glu Gln Val Arg Lys Asn Leu
 1090 1095 1100
 Asp Ile Ser Ile Asn Ala Ser Asn Asn Phe Asn Leu Asn Ile Thr Trp
 1105 1110 1115 1120
 Ser Val Gly Ser Ala Gly Thr Ile Ser Gly Glu Glu Thr Ser Ile Val
 1125 1130 1135
 Ser Lys Asn Asn Ile Lys Glu Tyr Arg Asp Ser Phe Ser Tyr Glu Lys
 1140 1145 1150
 Phe Asn Phe Arg Ser Asn Pro Asn Ile Thr Phe Tyr Val Tyr Val Ser
 1155 1160 1165

Asn Phe Ser Trp Pro Ile Lys Ile Gln Val Ser Val Lys Ser Ile Tyr
 1170 1175 1180

Phe
 1185

<210> 8
 <211> 6201
 <212> DNA
 <213> Homo sapiens

<400> 8

```

atgttgaagt tcaaatatgg agcgcggaat cctttggatg ctggtgctgc tgaacccatt 60
gccagccggg cctccaggct gaatctgttc ttccagggga aaccaccctt tatgactcaa 120
cagcagatgt ctcctctttc ccgagaaggg atattagatg ccctctttgt tctctttgaa 180
gaatgcagtc agcctgctct gatgaagatt aagcacgtga gcaactttgt ccggaagtgt 240
tccgacacca tagctgagtt acaggagctc cagccttcgg caaaggactt cgaagtcaga 300
agtcttgtag gttgtggtca ctttgcgtga gtgcaggtgg taagagagaa agcaaccggg 360
gacatctatg ctatgaaagt gatgaagaag aaggctttat tggcccagga gcaggtttca 420
ttttttgagg aagagcggaa catattatct cgaagcacia gcccgtggat cccccaatta 480
cagtatgcct ttcaggacaa aaatcacctt tatctggtga tggaaatca gcctggaggg 540
gacttgctgt cacttttgaa tagatatgag gaccagttag atgaaaacct gatacagttt 600
tacctagctg agctgatttt ggctgttcac agcgttcac t gatgggata cgtgcatcgg 660
gacatcaagc ctgagaacat tctcgttgac cgcacaggac acatcaagct ggtggatttt 720
ggatctgccg cgaaaatgaa ttcaaacaag gtgaatgcca aactcccgat tgggacccca 780
gattacatgg ctcctgaagt gctgactgtg atgaacgggg atggaaaagg cacctacggc 840
ctggactgtg actggtggtc agtgggcgtg attgcctatg agatgattta tgggagatcc 900
cccttcgcag agggaaacctc tgccagaacc ttcaataaca ttatgaattt ccagcggttt 960
ttgaaatttc cagatgaccc caaagtgagc agtgactttc ttgatctgat tcaaagcttg 1020
ttgtgcggcc agaaagagag actgaagttt gaaggctctt gctgccatcc tttcttctct 1080
aaaattgact ggaacaacat tcgtaacgct cctccccctc tcgttcccac cctcaagtct 1140
gacgatgaca cctccaattt tgatgaacca gagaagaatt cgtgggtttc atcctctccg 1200
tgccagctga gcccctcagg cttctcgggt gaagaactgc cgtttgtggg gttttcgtac 1260
agcaaggcac tggggattct tggtagatct gagtctgttg tgtcgggtct ggactcccc 1320
gccaagacta gctccatgga aaagaaactt ctcatcaaaa gcaaagagct acaagactct 1380
caggacaagt gtcacaagat ggagcaggaa atgacccggt tacatcggag agtgtcagag 1440
gtggaggctg tgcttagtca gaaggagggt gagctgaagg cctctgagac tcagagatcc 1500
ctcctggagc aggaccttgc tacctacatc acagaatgca gtagcttaaa gcgaagtttg 1560
gagcaagcac ggatggagggt gtcccaggag gatgacaaag cactgcagct tctccatgat 1620
atcagagagc agagccggaa gctccaagaa atcaaagagc aggagtacca ggctcaagtg 1680
gaagaaatga ggttgatgat gaatcagttg gaagaggatc ttgtctcagc aagaagacgg 1740
agtgatctct acgaatctga gctgagagag tctcggcttg ctgctgaaga attcaagcgg 1800
aaagcgacag aatgtcagca taaactgttg aaggctaagg atcaggggaa gcctgaagtg 1860
ggagaatatg cgaaactgga gaagatcaat gctgagcagc agctcaaaat tcaggagctc 1920
caagagaaac tggagaaggc tgtaaaagcc agcacggagg ccaccgagct gctgcagaat 1980
atccgccagg caaaggagcg agccgagagg gagctggaga agctgcagaa ccgagaggat 2040
tcttctgaag gcatcagaaa gaagctggtg gaagctgagg aacgccgcca ttctctggag 2100
aacaaggtaa agagactaga gaccatggag cgtagagaaa acagactgaa ggatgacatc 2160
cagacaaaat cccaacagat ccagcagatg gctgataaaa ttctggagct cgaagagaaa 2220
catcgggagg cccaagtctc agcccagcac ctagaagtgc acctgaaaca gaaagagcag 2280
cactatgagg aaaagattaa agtattggac aatcagataa agaaagacct ggctgacaag 2340
gagacactgg agaacatgat gcagagacac gaggaggagg cccatgagaa gggcaaaatt 2400
ctcagcgaac agaaggcgat gatcaatgct atggattcca agatcagatc cctggaacag 2460
aggattgtgg aactgtctga agccaataaa cttgcagcaa atagcagtct ttttacccaa 2520
aggaacatga aggcccaaga agagatgatt tctgaactca ggcaacagaa attttacctg 2580

```

gagacacagg	ctgggaagtt	ggaggcccag	aaccgaaaac	tggaggagca	gctggagaag	2640
atcagccacc	aagaccacag	tgacaagaat	cggctgctgg	aactggagac	aagattgctg	2700
gaggtgagtc	tagagcacga	ggagcagaaa	ctggagctca	agcgccagct	cacagagcta	2760
cagctctccc	tgcaggagcg	cgagtcacag	ttgacagccc	tgcaggctgc	acgggctggc	2820
ctggagagcc	agcttcgcca	ggcgaagaca	gagctggaag	agaccacagc	agaagctgaa	2880
gaggagatcc	aggcactcac	ggcacataga	gatgaaatcc	agcgcaaatt	tgatgctctt	2940
cgtaacagct	gtactgtgat	cacagacctg	gaggagcagc	taaaccagct	gaccgaggac	3000
aacgctgaac	tcaacaacca	aaactttctac	ttgtccaaac	aactcgatga	ggcttctggc	3060
gccaacgacg	agattgtaca	actgcgaagt	gaagtggacc	atctccgccc	ggagatcacg	3120
gaacgagaga	tgcagcttac	cagccagaag	caaacgatgg	aggctctgaa	gaccacgtgc	3180
accatgctgg	aggaacaggt	catggatttg	gaggccctaa	acgatgagct	gctagaaaaa	3240
gagcggcagt	gggaggcctg	gaggagcgct	ctgggtgatg	agaaatccca	gtttgagtgt	3300
cgggttcgag	agctgcagag	gatgctggac	accgagaaaac	agagcagggc	gagagccgat	3360
cagcggatca	ccgagctctg	ccagggtggt	gagctggcag	tgaaggagca	caaggctgag	3420
attctcgctc	tgcagcaggc	tctcaaagag	cagaagctga	aggccgagag	cctctctgac	3480
aagctcaatg	acctggagaa	gaagcatgct	atgcttgaaa	tgaatgcccg	aagcttacag	3540
cagaagctgg	agactgaacg	agagctcaaa	cagaggcttc	tggaagagca	agccaaatta	3600
cagcagcaga	tggacctgca	gaaaaatcac	attttccgct	tgactcaagg	actgcaagaa	3660
gctctagatc	gggctgatct	actgaagaca	gaaagaagtg	acttgagta	tcagctggaa	3720
aacattcagg	tgctctatct	tcatgaaaag	gtgaaaatgg	aaggcactat	ttctcaacaa	3780
accaaactca	ttgattttct	gcaagccaaa	atggaccaac	ctgctaaaaa	gaaaaagggtg	3840
cctctgcagt	acaatgagct	gaagctggcc	ctggagaagg	agaaagctcg	ctgtgcagag	3900
ctagaggaag	cccttcagaa	gacccgcctc	gagctccggt	ccgcccggga	ggaagctgcc	3960
caccgcaaag	caacggacca	cccacaccca	tccacgccag	ccaccgcgag	gcagcagatc	4020
gccatgtctg	ccatcgctgc	gtcgccagag	caccagccca	gtgccatgag	cctgctggcc	4080
ccgccatcca	gccgcagaaa	ggagtcttca	actccagagg	aatttagtgc	gcgtcttaag	4140
gaacgcatgc	accacaatat	tcttcaccga	ttcaacgtag	gactgaacat	gcgagccaca	4200
aagtgtgctg	tgtgtctgga	taccgtgcac	tttgacgcc	aggcatccaa	atgtctagaa	4260
tgctcaggtg	tgtgtcaccc	caagtgtctc	acgtgcttgc	cagccacctg	cggcttgctc	4320
gctgaatatg	ccacacactt	caccgaggcc	ttctgccgtg	acaaaatgaa	ctccccaggt	4380
ctccagacca	aggagcccag	cagcagcttg	cacctggaag	ggtggatgaa	ggtgcccagg	4440
aataacaaac	gaggacagca	aggctgggac	aggaagtaca	ttgtcctgga	gggatcaaaa	4500
gtcctcattt	atgacaatga	agccagagaa	gctggacaga	ggccggtgga	agaatttgag	4560
ctgtgccttc	ccgacgggga	tgtatctatt	catggtgccg	ttggtgcttc	cgaactcgca	4620
aatacagcca	aagcagatgt	cccatacata	ctgaagatgg	aatctcacc	gcacaccacc	4680
tgctggcccc	ggagaaccct	ctacttgcta	gctcccagct	tccctgacaa	acagcgctgg	4740
gtcaccgcct	tagaatcagt	tgtcgcaggt	gggagagttt	ctagggaaaa	agcagaagct	4800
gatgctaaac	tgcttggaag	ctccctgctg	aaactggaag	gtgatgaccg	tctagacatg	4860
aactgcacgc	tgcccttcag	tgaccaggtg	gtgttggtgg	gcaccgagga	agggtcttac	4920
gccctgaatg	tcttgaaaaa	ctccctaacc	catgtcccag	gaattggagc	agtcttccaa	4980
atttatatta	tcaaggacct	ggagaagcta	ctcatgatag	caggtgaaga	gcgggcaactg	5040
tgtcttggtg	acgtgaagaa	agtgaacag	tccctggccc	agtcccacct	gctgcccag	5100
ccgacatct	cacccaacat	ttttgaagct	gtcaagggct	gccacttggt	tggggcaggc	5160
aagattgaga	acgggctctg	catctgtgca	gccatgcccc	gcaaagtcgt	cattctccgc	5220
tacaacgaaa	acctcagcaa	atactgcatc	cggaaagaga	tagagacctc	agagccctgc	5280
agctgtatcc	acttcaccaa	ttacagtatc	ctcattggaa	ccaataaatt	ctacgaaatc	5340
gacatgaagc	agtacacgct	cgaggaattc	ctggataaga	atgaccattc	cttggcacct	5400
gctgtgtttg	ccgcctcttc	caacagcttc	cctgtctcaa	tcgtgcaggt	gaacagcgca	5460
gggcagcgag	aggagtactt	gctgtgtttc	cacgaatttg	gagtgttcgt	ggattcttac	5520
ggaagacgta	gccgcacaga	cgatctcaag	tggagtgcgt	tacctttggc	ctttgcctac	5580
agagaaccct	atctgtttgt	gacccacttc	aactcactcg	aagtaattga	gatccaggca	5640
cgctcctcag	cagggacccc	tgcccagagc	tacctggaca	tcccgaacct	gcgctacctg	5700
ggccttgcca	tttctctcag	agcgatttac	ttggcgctct	cataccagga	taaattaagg	5760
gtcatttgct	gcaagggaaa	cctcgtgaag	gagtccggca	ctgaacacca	ccggggcccc	5820
tccacctccc	gcagcagccc	caacaagcga	ggcccaccca	cgtacaacga	gcacatcacc	5880
aagcgcgtgg	cctccagccc	agcgcgcccc	gaaggcccca	gccaccgcg	agagccaagc	5940
acaccccacc	gctaccgcga	ggggcggacc	gagctgcgca	gggacaagtc	tcctggccgc	6000

```

ccccctggagc gagagaagtc ccccgcccg atgctcagca cgcggagaga gcggtcccc 6060
gggaggctgt ttgaagacag cagcaggggc cggctgcctg cgggagccgt gaggacccc 6120
ctgtcccagg tgaacaaggt gaggcagcat tccgaggcct gtgtgtctgt tgcggaggcc 6180
aggagtgact tggggaactg a                                     6201

```

```

<210> 9
<211> 2066
<212> PRT
<213> Homo sapiens

```

```

<400> 9
Met Leu Lys Phe Lys Tyr Gly Ala Arg Asn Pro Leu Asp Ala Gly Ala
 1             5             10             15

Ala Glu Pro Ile Ala Ser Arg Ala Ser Arg Leu Asn Leu Phe Phe Gln
      20             25             30

Gly Lys Pro Pro Phe Met Thr Gln Gln Gln Met Ser Pro Leu Ser Arg
      35             40             45

Glu Gly Ile Leu Asp Ala Leu Phe Val Leu Phe Glu Glu Cys Ser Gln
      50             55             60

Pro Ala Leu Met Lys Ile Lys His Val Ser Asn Phe Val Arg Lys Cys
      65             70             75             80

Ser Asp Thr Ile Ala Glu Leu Gln Glu Leu Gln Pro Ser Ala Lys Asp
      85             90             95

Phe Glu Val Arg Ser Leu Val Gly Cys Gly His Phe Ala Glu Val Gln
      100            105            110

Val Val Arg Glu Lys Ala Thr Gly Asp Ile Tyr Ala Met Lys Val Met
      115            120            125

Lys Lys Lys Ala Leu Leu Ala Gln Glu Gln Val Ser Phe Phe Glu Glu
      130            135            140

Glu Arg Asn Ile Leu Ser Arg Ser Thr Ser Pro Trp Ile Pro Gln Leu
      145            150            155            160

Gln Tyr Ala Phe Gln Asp Lys Asn His Leu Tyr Leu Val Met Glu Tyr
      165            170            175

Gln Pro Gly Gly Asp Leu Leu Ser Leu Leu Asn Arg Tyr Glu Asp Gln
      180            185            190

Leu Asp Glu Asn Leu Ile Gln Phe Tyr Leu Ala Glu Leu Ile Leu Ala
      195            200            205

Val His Ser Val His Leu Met Gly Tyr Val His Arg Asp Ile Lys Pro
      210            215            220

Glu Asn Ile Leu Val Asp Arg Thr Gly His Ile Lys Leu Val Asp Phe
      225            230            235            240

```

Gly Ser Ala Ala Lys Met Asn Ser Asn Lys Val Asn Ala Lys Leu Pro
 245 250 255
 Ile Gly Thr Pro Asp Tyr Met Ala Pro Glu Val Leu Thr Val Met Asn
 260 265 270
 Gly Asp Gly Lys Gly Thr Tyr Gly Leu Asp Cys Asp Trp Trp Ser Val
 275 280 285
 Gly Val Ile Ala Tyr Glu Met Ile Tyr Gly Arg Ser Pro Phe Ala Glu
 290 295 300
 Gly Thr Ser Ala Arg Thr Phe Asn Asn Ile Met Asn Phe Gln Arg Phe
 305 310 315 320
 Leu Lys Phe Pro Asp Asp Pro Lys Val Ser Ser Asp Phe Leu Asp Leu
 325 330 335
 Ile Gln Ser Leu Leu Cys Gly Gln Lys Glu Arg Leu Lys Phe Glu Gly
 340 345 350
 Leu Cys Cys His Pro Phe Phe Ser Lys Ile Asp Trp Asn Asn Ile Arg
 355 360 365
 Asn Ala Pro Pro Pro Phe Val Pro Thr Leu Lys Ser Asp Asp Asp Thr
 370 375 380
 Ser Asn Phe Asp Glu Pro Glu Lys Asn Ser Trp Val Ser Ser Ser Pro
 385 390 395 400
 Cys Gln Leu Ser Pro Ser Gly Phe Ser Gly Glu Glu Leu Pro Phe Val
 405 410 415
 Gly Phe Ser Tyr Ser Lys Ala Leu Gly Ile Leu Gly Arg Ser Glu Ser
 420 425 430
 Val Val Ser Gly Leu Asp Ser Pro Ala Lys Thr Ser Ser Met Glu Lys
 435 440 445
 Lys Leu Leu Ile Lys Ser Lys Glu Leu Gln Asp Ser Gln Asp Lys Cys
 450 455 460
 His Lys Met Glu Gln Glu Met Thr Arg Leu His Arg Arg Val Ser Glu
 465 470 475 480
 Val Glu Ala Val Leu Ser Gln Lys Glu Val Glu Leu Lys Ala Ser Glu
 485 490 495
 Thr Gln Arg Ser Leu Leu Glu Gln Asp Leu Ala Thr Tyr Ile Thr Glu
 500 505 510
 Cys Ser Ser Leu Lys Arg Ser Leu Glu Gln Ala Arg Met Glu Val Ser
 515 520 525
 Gln Glu Asp Asp Lys Ala Leu Gln Leu Leu His Asp Ile Arg Glu Gln
 530 535 540

Ser Arg Lys Leu Gln Glu Ile Lys Glu Gln Glu Tyr Gln Ala Gln Val
 545 550 555 560
 Glu Glu Met Arg Leu Met Met Asn Gln Leu Glu Glu Asp Leu Val Ser
 565 570 575
 Ala Arg Arg Arg Ser Asp Leu Tyr Glu Ser Glu Leu Arg Glu Ser Arg
 580 585 590
 Leu Ala Ala Glu Glu Phe Lys Arg Lys Ala Thr Glu Cys Gln His Lys
 595 600 605
 Leu Leu Lys Ala Lys Asp Gln Gly Lys Pro Glu Val Gly Glu Tyr Ala
 610 615 620
 Lys Leu Glu Lys Ile Asn Ala Glu Gln Gln Leu Lys Ile Gln Glu Leu
 625 630 635 640
 Gln Glu Lys Leu Glu Lys Ala Val Lys Ala Ser Thr Glu Ala Thr Glu
 645 650 655
 Leu Leu Gln Asn Ile Arg Gln Ala Lys Glu Arg Ala Glu Arg Glu Leu
 660 665 670
 Glu Lys Leu Gln Asn Arg Glu Asp Ser Ser Glu Gly Ile Arg Lys Lys
 675 680 685
 Leu Val Glu Ala Glu Glu Arg Arg His Ser Leu Glu Asn Lys Val Lys
 690 695 700
 Arg Leu Glu Thr Met Glu Arg Arg Glu Asn Arg Leu Lys Asp Asp Ile
 705 710 715 720
 Gln Thr Lys Ser Gln Gln Ile Gln Gln Met Ala Asp Lys Ile Leu Glu
 725 730 735
 Leu Glu Glu Lys His Arg Glu Ala Gln Val Ser Ala Gln His Leu Glu
 740 745 750
 Val His Leu Lys Gln Lys Glu Gln His Tyr Glu Glu Lys Ile Lys Val
 755 760 765
 Leu Asp Asn Gln Ile Lys Lys Asp Leu Ala Asp Lys Glu Thr Leu Glu
 770 775 780
 Asn Met Met Gln Arg His Glu Glu Glu Ala His Glu Lys Gly Lys Ile
 785 790 795 800
 Leu Ser Glu Gln Lys Ala Met Ile Asn Ala Met Asp Ser Lys Ile Arg
 805 810 815
 Ser Leu Glu Gln Arg Ile Val Glu Leu Ser Glu Ala Asn Lys Leu Ala
 820 825 830
 Ala Asn Ser Ser Leu Phe Thr Gln Arg Asn Met Lys Ala Gln Glu Glu
 835 840 845

Met Ile Ser Glu Leu Arg Gln Gln Lys Phe Tyr Leu Glu Thr Gln Ala
 850 855 860
 Gly Lys Leu Glu Ala Gln Asn Arg Lys Leu Glu Glu Gln Leu Glu Lys
 865 870 875 880
 Ile Ser His Gln Asp His Ser Asp Lys Asn Arg Leu Leu Glu Leu Glu
 885 890 895
 Thr Arg Leu Arg Glu Val Ser Leu Glu His Glu Glu Gln Lys Leu Glu
 900 905 910
 Leu Lys Arg Gln Leu Thr Glu Leu Gln Leu Ser Leu Gln Glu Arg Glu
 915 920 925
 Ser Gln Leu Thr Ala Leu Gln Ala Ala Arg Ala Ala Leu Glu Ser Gln
 930 935 940
 Leu Arg Gln Ala Lys Thr Glu Leu Glu Glu Thr Thr Ala Glu Ala Glu
 945 950 955 960
 Glu Glu Ile Gln Ala Leu Thr Ala His Arg Asp Glu Ile Gln Arg Lys
 965 970 975
 Phe Asp Ala Leu Arg Asn Ser Cys Thr Val Ile Thr Asp Leu Glu Glu
 980 985 990
 Gln Leu Asn Gln Leu Thr Glu Asp Asn Ala Glu Leu Asn Asn Gln Asn
 995 1000 1005
 Phe Tyr Leu Ser Lys Gln Leu Asp Glu Ala Ser Gly Ala Asn Asp Glu
 1010 1015 1020
 Ile Val Gln Leu Arg Ser Glu Val Asp His Leu Arg Arg Glu Ile Thr
 1025 1030 1035 1040
 Glu Arg Glu Met Gln Leu Thr Ser Gln Lys Gln Thr Met Glu Ala Leu
 1045 1050 1055
 Lys Thr Thr Cys Thr Met Leu Glu Glu Gln Val Met Asp Leu Glu Ala
 1060 1065 1070
 Leu Asn Asp Glu Leu Leu Glu Lys Glu Arg Gln Trp Glu Ala Trp Arg
 1075 1080 1085
 Ser Val Leu Gly Asp Glu Lys Ser Gln Phe Glu Cys Arg Val Arg Glu
 1090 1095 1100
 Leu Gln Arg Met Leu Asp Thr Glu Lys Gln Ser Arg Ala Arg Ala Asp
 1105 1110 1115 1120
 Gln Arg Ile Thr Glu Ser Arg Gln Val Val Glu Leu Ala Val Lys Glu
 1125 1130 1135
 His Lys Ala Glu Ile Leu Ala Leu Gln Gln Ala Leu Lys Glu Gln Lys
 1140 1145 1150

Leu Lys Ala Glu Ser Leu Ser Asp Lys Leu Asn Asp Leu Glu Lys Lys
 1155 1160 1165
 His Ala Met Leu Glu Met Asn Ala Arg Ser Leu Gln Gln Lys Leu Glu
 1170 1175 1180
 Thr Glu Arg Glu Leu Lys Gln Arg Leu Leu Glu Glu Gln Ala Lys Leu
 1185 1190 1195 1200
 Gln Gln Gln Met Asp Leu Gln Lys Asn His Ile Phe Arg Leu Thr Gln
 1205 1210 1215
 Gly Leu Gln Glu Ala Leu Asp Arg Ala Asp Leu Leu Lys Thr Glu Arg
 1220 1225 1230
 Ser Asp Leu Glu Tyr Gln Leu Glu Asn Ile Gln Val Leu Tyr Ser His
 1235 1240 1245
 Glu Lys Val Lys Met Glu Gly Thr Ile Ser Gln Gln Thr Lys Leu Ile
 1250 1255 1260
 Asp Phe Leu Gln Ala Lys Met Asp Gln Pro Ala Lys Lys Lys Lys Val
 1265 1270 1275 1280
 Pro Leu Gln Tyr Asn Glu Leu Lys Leu Ala Leu Glu Lys Glu Lys Ala
 1285 1290 1295
 Arg Cys Ala Glu Leu Glu Glu Ala Leu Gln Lys Thr Arg Ile Glu Leu
 1300 1305 1310
 Arg Ser Ala Arg Glu Glu Ala Ala His Arg Lys Ala Thr Asp His Pro
 1315 1320 1325
 His Pro Ser Thr Pro Ala Thr Ala Arg Gln Gln Ile Ala Met Ser Ala
 1330 1335 1340
 Ile Val Arg Ser Pro Glu His Gln Pro Ser Ala Met Ser Leu Leu Ala
 1345 1350 1355 1360
 Pro Pro Ser Ser Arg Arg Lys Glu Ser Ser Thr Pro Glu Glu Phe Ser
 1365 1370 1375
 Arg Arg Leu Lys Glu Arg Met His His Asn Ile Pro His Arg Phe Asn
 1380 1385 1390
 Val Gly Leu Asn Met Arg Ala Thr Lys Cys Ala Val Cys Leu Asp Thr
 1395 1400 1405
 Val His Phe Gly Arg Gln Ala Ser Lys Cys Leu Glu Cys Gln Val Met
 1410 1415 1420
 Cys His Pro Lys Cys Ser Thr Cys Leu Pro Ala Thr Cys Gly Leu Pro
 1425 1430 1435 1440
 Ala Glu Tyr Ala Thr His Phe Thr Glu Ala Phe Cys Arg Asp Lys Met
 1445 1450 1455

Asn Ser Pro Gly Leu Gln Thr Lys Glu Pro Ser Ser Ser Leu His Leu
 1460 1465 1470
 Glu Gly Trp Met Lys Val Pro Arg Asn Asn Lys Arg Gly Gln Gln Gly
 1475 1480 1485
 Trp Asp Arg Lys Tyr Ile Val Leu Glu Gly Ser Lys Val Leu Ile Tyr
 1490 1495 1500
 Asp Asn Glu Ala Arg Glu Ala Gly Gln Arg Pro Val Glu Glu Phe Glu
 1505 1510 1515 1520
 Leu Cys Leu Pro Asp Gly Asp Val Ser Ile His Gly Ala Val Gly Ala
 1525 1530 1535
 Ser Glu Leu Ala Asn Thr Ala Lys Ala Asp Val Pro Tyr Ile Leu Lys
 1540 1545 1550
 Met Glu Ser His Pro His Thr Thr Cys Trp Pro Gly Arg Thr Leu Tyr
 1555 1560 1565
 Leu Leu Ala Pro Ser Phe Pro Asp Lys Gln Arg Trp Val Thr Ala Leu
 1570 1575 1580
 Glu Ser Val Val Ala Gly Gly Arg Val Ser Arg Glu Lys Ala Glu Ala
 1585 1590 1595 1600
 Asp Ala Lys Leu Leu Gly Asn Ser Leu Leu Lys Leu Glu Gly Asp Asp
 1605 1610 1615
 Arg Leu Asp Met Asn Cys Thr Leu Pro Phe Ser Asp Gln Val Val Leu
 1620 1625 1630
 Val Gly Thr Glu Glu Gly Leu Tyr Ala Leu Asn Val Leu Lys Asn Ser
 1635 1640 1645
 Leu Thr His Val Pro Gly Ile Gly Ala Val Phe Gln Ile Tyr Ile Ile
 1650 1655 1660
 Lys Asp Leu Glu Lys Leu Leu Met Ile Ala Gly Glu Glu Arg Ala Leu
 1665 1670 1675 1680
 Cys Leu Val Asp Val Lys Lys Val Lys Gln Ser Leu Ala Gln Ser His
 1685 1690 1695
 Leu Pro Ala Gln Pro Asp Ile Ser Pro Asn Ile Phe Glu Ala Val Lys
 1700 1705 1710
 Gly Cys His Leu Phe Gly Ala Gly Lys Ile Glu Asn Gly Leu Cys Ile
 1715 1720 1725
 Cys Ala Ala Met Pro Ser Lys Val Val Ile Leu Arg Tyr Asn Glu Asn
 1730 1735 1740
 Leu Ser Lys Tyr Cys Ile Arg Lys Glu Ile Glu Thr Ser Glu Pro Cys
 1745 1750 1755 1760

Ser Cys Ile His Phe Thr Asn Tyr Ser Ile Leu Ile Gly Thr Asn Lys
 1765 1770 1775
 Phe Tyr Glu Ile Asp Met Lys Gln Tyr Thr Leu Glu Glu Phe Leu Asp
 1780 1785 1790
 Lys Asn Asp His Ser Leu Ala Pro Ala Val Phe Ala Ala Ser Ser Asn
 1795 1800 1805
 Ser Phe Pro Val Ser Ile Val Gln Val Asn Ser Ala Gly Gln Arg Glu
 1810 1815 1820
 Glu Tyr Leu Leu Cys Phe His Glu Phe Gly Val Phe Val Asp Ser Tyr
 1825 1830 1835 1840
 Gly Arg Arg Ser Arg Thr Asp Asp Leu Lys Trp Ser Arg Leu Pro Leu
 1845 1850 1855
 Ala Phe Ala Tyr Arg Glu Pro Tyr Leu Phe Val Thr His Phe Asn Ser
 1860 1865 1870
 Leu Glu Val Ile Glu Ile Gln Ala Arg Ser Ser Ala Gly Thr Pro Ala
 1875 1880 1885
 Arg Ala Tyr Leu Asp Ile Pro Asn Pro Arg Tyr Leu Gly Pro Ala Ile
 1890 1895 1900
 Ser Ser Gly Ala Ile Tyr Leu Ala Ser Ser Tyr Gln Asp Lys Leu Arg
 1905 1910 1915 1920
 Val Ile Cys Cys Lys Gly Asn Leu Val Lys Glu Ser Gly Thr Glu His
 1925 1930 1935
 His Arg Gly Pro Ser Thr Ser Arg Ser Ser Pro Asn Lys Arg Gly Pro
 1940 1945 1950
 Pro Thr Tyr Asn Glu His Ile Thr Lys Arg Val Ala Ser Ser Pro Ala
 1955 1960 1965
 Pro Pro Glu Gly Pro Ser His Pro Arg Glu Pro Ser Thr Pro His Arg
 1970 1975 1980
 Tyr Arg Glu Gly Arg Thr Glu Leu Arg Arg Asp Lys Ser Pro Gly Arg
 1985 1990 1995 2000
 Pro Leu Glu Arg Glu Lys Ser Pro Gly Arg Met Leu Ser Thr Arg Arg
 2005 2010 2015
 Glu Arg Ser Pro Gly Arg Leu Phe Glu Asp Ser Ser Arg Gly Arg Leu
 2020 2025 2030
 Pro Ala Gly Ala Val Arg Thr Pro Leu Ser Gln Val Asn Lys Val Arg
 2035 2040 2045
 Gln His Ser Glu Ala Cys Val Ser Val Ala Glu Ala Arg Ser Asp Leu
 2050 2055 2060

Gly Asn
2065

<210> 10
<211> 6189
<212> DNA
<213> Homo sapiens

<400> 10

```
atgttgaagt tcaaatatgg agcgcggaat cctttggatg ctggtgctgc tgaacccatt 60
gccagccggg cctccaggct gaatctgttc ttccagggga aaccaccctt tatgactcaa 120
cagcagatgt ctctcttttc ccgagaaggg atattagatg ccctctttgt tctctttgaa 180
gaatgcagtc agcctgctct gatgaagatt aagcacgtga gcaactttgt ccggaagtgt 240
tccgacacca tagctgagtt acaggagctc cagccttcgg caaaggactt cgaagtcaga 300
agtctttag tagttgggtca ctttgctgaa gtgcagggtg taagagagaa agcaaccggg 360
gacatctatg ctatgaaagt gatgaagaag aaggctttat tggcccagga gcaggtttca 420
ttttttgagg aagagcggaa catattatct cgaagcacia gccctggat cccccaatta 480
cagtatgcct ttcaggacaa aaatcacctt tatctggtga tggaatatca gcctggaggg 540
gacttgctgt cacttttgaa tagatatgag gaccagttag atgaaaacct gatacagttt 600
tacctagctg agctgatttt ggctgttcac agcgttcac tgatgggata cgtgcacg 660
gacatcaagc ctgagaacat tctcgttgac cgcacaggac acatcaagct ggtggatttt 720
ggatctgccg cgaaaatgaa ttcaaacaag gtgaatgcca aactcccgat tgggacccca 780
gattacatgg ctctgaagt gctgactgtg atgaacgggg atggaaaagg cacctacggc 840
ctggactgtg actggtggtc agtgggcgtg attgcctatg agatgattta tgggagatcc 900
cccttcgcag agggaacctc tgccagaacc ttcaataaca ttatgaattt ccagcggttt 960
ttgaaatttc cagatgacct caaagtgagc agtgactttc ttgatctgat tcaaagcttg 1020
ttgtgcggcc agaaagagag actgaagttt gaaggtcttt gctgccatcc tttctctct 1080
aaaattgact ggaacaacat tcgtaacgct cctccccctc tcgttccac cctcaagtct 1140
gacgatgaca cctccaattt tgatgaacca gagaagaatt cgtgggtttc atcctctccg 1200
tgccagctga gcccctcagg cttctcgggt gaagaactgc cgtttgtggg gtttctgtac 1260
agcaaggcac tggggattct tggtagatct gagtctgttg tgcgggtct ggactccct 1320
gccaagacta gctccatgga aaagaaactt ctcacaaaaa gcaaagagct acaagactct 1380
caggacaagt gtcacaagat ggagcaggaa atgaccgggt tacatcggag agtgtcagag 1440
gtggaggctg tgcttagtca gaaggagtg gagctgaagg cctctgagac tcagagatcc 1500
ctcttgagc aggacctgc tacctacac acagaatgca gtagcttaa gcaagtttg 1560
gagcaagcac ggatggaggt gtcccaggag gatgacaaa cactgcagct tctccatgat 1620
atcagagagc agagccggaa gctccaagaa atcaaagagc aggagtacca gggtcaagt 1680
gaagaaatga ggttgatgat gaatcagttg gaagaggatc ttgtctcagc aagaagacgg 1740
agtgatctct acgaatctga gctgagagag tctcggcttg ctgctgaaga attcaagcgg 1800
aaagcgacag aatgtcagca taaactgttg aaggctaagg atcaggggaa gcctgaagt 1860
ggagaatatg cgaaaactgga gaagatcaat gctgagcagc agtcaaaaat tcaggagctc 1920
caagagaaac tggagaaggc tgtaaaagcc agcagggagg ccaccgagct gctgcagaat 1980
atccgccagg caaaggagcg agccgagagg gagctggaga agctgcagaa ccgagaggat 2040
tcttctgaag gcatcagaaa gaagctggtg gaagctgagg aacgccgcca ttctctggag 2100
aacaaggtaa agagactaga gaccatggag cgtagagaaa acagactgaa ggatgacatc 2160
cagacaaaat cccaacagat ccagcagatg gctgataaaa ttctggagct cgaagagaaa 2220
catcgggagg cccaagtctc agcccagcac ctagaagtgc acctgaaaca gaaagagcag 2280
cactatgagg aaaagattaa agtattggac aatcagataa agaaagacct ggctgacaag 2340
gagacactgg agaacatgat gcagagacac gaggaggagg ccatgagaa gggcaaaatt 2400
ctcagcgaac agaaggcgat gatcaatgct atggattcca agatcagatc cctggaacag 2460
aggattgtgg aactgtctga agccaataaa cttgcagcaa atagcagtct ttttacccaa 2520
aggaacatga aggccaaga agagatgatt tctgaactca ggcaacagaa attttacctg 2580
gagacacagg ctgggaagtt ggaggcccag aaccgaaaac tggaggagca gctggagaag 2640
atcagccacc aagaccacag tgacaagaat cggctgctgg aactggagac aagattgcgg 2700
gaggtgagtc tagagcacga ggagcagaaa ctggagctca agcgcagct cacagagcta 2760
cagctctccc tgcaggagcg cgagtcacag ttgacagccc tgcaggctgc acgggcggcc 2820
```

ctggagagcc	agcttcgcca	ggcgaagaca	gagctggaag	agaccacagc	agaagctgaa	2880
gaggagatcc	aggcactcac	ggcacataga	gatgaaatcc	agcgcaaatt	tgatgctctt	2940
cgtaacagct	gtactgtgat	cacagacctg	gaggagcagc	taaaccagct	gaccgaggac	3000
aacgctgaac	tcaacaacca	aaactttctac	ttgtccaaac	aactcgatga	ggctttctggc	3060
gccaacgacg	agattgtaca	actgcgaagt	gaagtggacc	atctccgccg	ggagatcacg	3120
gaacgagaga	tgcagcttac	cagccagaag	caaacgatgg	aggctctgaa	gaccacgtgc	3180
accatgctgg	aggaacaggt	catggatttg	gaggccctaa	acgatgagct	gctagaaaaa	3240
gagcggcagt	gggaggcctg	gaggagcgtc	ctgggtgatg	agaaatccca	gtttgagtgt	3300
cgggttcgag	agctgcagag	gatgctggac	accgagaaac	agagcagggc	gagagccgat	3360
cagcggatca	ccgagtctcg	ccagggtggtg	gagctggcag	tgaaggagca	caaggctgag	3420
attctcgctc	tgcagcaggc	tctcaaagag	cagaagctga	aggccgagag	cctctctgac	3480
aagctcaatg	acctggagaa	gaagcatgct	atgcttgaaa	tgaatgcccg	aagcttacag	3540
cagaagctgg	agactgaacg	agagctcaaa	cagaggcttc	tggaagagca	agccaaatta	3600
cagcagcaga	tggacctgca	gaaaaatcac	attttccgtc	tgactcaagg	actgcaagaa	3660
gctctagatc	gggctgatct	actgaagaca	gaaagaagtg	acttgagta	tcagctggaa	3720
aacattcagg	tgctctattc	tcatgaaaag	gtgaaaatgg	aaggcactat	ttctcaacaa	3780
accaaactca	ttgattttct	gcaagccaaa	atggaccaac	ctgctaaaaa	gaaaaaggtg	3840
cctctgcagt	acaatgagct	gaagctggcc	ctggagaagg	agaaagctcg	ctgtgcagag	3900
ctagaggaag	cccttcagaa	gacccgcctc	gagctccggt	ccgcccggga	ggaagctgcc	3960
caccgcaaag	caacggacca	cccacaccca	tccacgcag	ccaccgcgag	gcagcagatc	4020
gccatgtctg	ccatcgtgcg	gtcgccagag	caccagccca	gtgccatgag	cctgctggcc	4080
ccgccatcca	gccgcagaaa	ggagtcttca	actccagagg	aatttagtcg	gcgtcttaag	4140
gaacgcatgc	accacaatat	tcctcaccga	ttcaacgtag	gactgaacat	gcgagccaca	4200
aagtgtgctg	tgtgtctgga	taccgtgcac	tttgagcgcc	aggcatccaa	atgtctagaa	4260
tgtcagggtga	tgtgtcacc	caagtgtctc	acgtgcttgc	cagccacctg	cggcttgccct	4320
gctgaatatg	ccacacactt	caccgaggcc	ttctgcccgtg	acaaaatgaa	ctccccaggt	4380
ctccagacca	aggagcccag	cagcagcttg	cacctggaag	ggtggatgaa	ggtgcccagg	4440
aataacaaac	gaggacagca	aggctgggac	aggaagtaca	ttgtcctgga	gggatcaaaa	4500
gtcctcattt	atgacaatga	agccagagaa	gctggacaga	ggccggtgga	agaatttgag	4560
ctgtgccttc	ccgacgggga	tgtatctatt	catggtgccg	ttggtgcttc	cgaactcgca	4620
aatacagcca	aagcagatgt	cccatacata	ctgaagatgg	aatctcacc	gcacaccacc	4680
tgctggcccc	ggagaaccct	ctacttgcta	gctcccagct	tccctgacaa	acagcgctgg	4740
gtcaccgcct	tagaatcagt	tgtcgcaggt	gggagagttt	ctagggaaaa	agcagaagct	4800
gatgctaaac	tgcttggaag	ctccctgctg	aaactggaag	gtgatgaccg	tctagacatg	4860
aactgcacgc	tgcccttcag	tgaccaggta	gtgttggtgg	gcaccgagga	agggtcttac	4920
gcctgaatg	tcttgaaaaa	ctccctaacc	catgtcccag	gaattggagc	agtcttccaa	4980
atttatatta	tcaaggacct	ggagaagcta	ctcatgatag	cagggtgaaga	gcgggcaactg	5040
tgtcttggtg	acgtgaagaa	agtgaacag	tccctggccc	agtcccacct	gcctgcccag	5100
cccagacatc	caccacaacat	ttttgaagct	gtcaagggtc	gccacttggt	tggggcaggc	5160
agatttga	acgggctctg	catctgtgca	gccatgcccc	gcaaagtcgt	cattctccgc	5220
tacaacgaaa	acctcagcaa	atactgcac	cggaaagaga	tagagacctc	agagccctgc	5280
agctgtatcc	acttcaccaa	ttacagtatc	ctcattggaa	ccaataaatt	ctacgaaatc	5340
gacatgaagc	agtacacgct	cgaggaattc	ctggataaga	atgaccattc	cttggcacct	5400
gctgtgtttg	ccgcctcttc	caacagcttc	cctgtctcaa	tcgtgcaggt	gaacagcgca	5460
gggcagcgag	aggagtactt	gctgtgtttc	cacgaatttg	gagtgttcgt	ggattcttac	5520
ggaagacgta	gccgcacaga	cgatctcaag	tggagtgcgt	tacctttggc	ctttgcctac	5580
agagaaccct	atctgtttgt	gacccaactc	aactcactcg	aagtaattga	gatccaggca	5640
cgctcctcag	cagggacccc	tgcccagagc	tacctggaca	tcccgaaccc	gcgctacctg	5700
ggccctgcc	tttctcagg	agcgatttac	ttggcgtcct	cataccagga	taaattaagg	5760
gtcatttgct	gcaagggaaa	cctcgtgaag	gagtccggca	ctgaacacca	ccggggcccc	5820
tccacctccc	gcagcagccc	caacaagcga	ggcccaccca	cgtacaacga	gcacatcacc	5880
aagcgcgtgg	cctccagccc	agcgcgcgcc	gaaggcccca	gccaccgcg	agagccaagc	5940
acacccacc	gctaccgcga	ggggcggacc	gagctgcgca	gggacaagtc	tcctggccgc	6000
cccctggagc	gagagaagtc	ccccggccgg	atgctcagca	cgcgagagga	gcggtcccc	6060
gggaggctgt	ttgaagacag	cagcaggggc	cggctgcctg	cgggagccgt	gaggaccccc	6120
ctgtcccagg	tgaacaaggt	gtgggaccag	tcttcagtat	aaatctcagc	cagaaaaaac	6180
aactcctca						6189

<210> 11
 <211> 2053
 <212> PRT
 <213> Homo sapiens

<400> 11

Met	Leu	Lys	Phe	Lys	Tyr	Gly	Ala	Arg	Asn	Pro	Leu	Asp	Ala	Gly	Ala
1				5					10					15	
Ala	Glu	Pro	Ile	Ala	Ser	Arg	Ala	Ser	Arg	Leu	Asn	Leu	Phe	Phe	Gln
			20					25					30		
Gly	Lys	Pro	Pro	Phe	Met	Thr	Gln	Gln	Gln	Met	Ser	Pro	Leu	Ser	Arg
		35					40					45			
Glu	Gly	Ile	Leu	Asp	Ala	Leu	Phe	Val	Leu	Phe	Glu	Glu	Cys	Ser	Gln
	50					55					60				
Pro	Ala	Leu	Met	Lys	Ile	Lys	His	Val	Ser	Asn	Phe	Val	Arg	Lys	Cys
65					70					75					80
Ser	Asp	Thr	Ile	Ala	Glu	Leu	Gln	Glu	Leu	Gln	Pro	Ser	Ala	Lys	Asp
			85						90					95	
Phe	Glu	Val	Arg	Ser	Leu	Val	Gly	Cys	Gly	His	Phe	Ala	Glu	Val	Gln
			100					105					110		
Val	Val	Arg	Glu	Lys	Ala	Thr	Gly	Asp	Ile	Tyr	Ala	Met	Lys	Val	Met
		115					120					125			
Lys	Lys	Lys	Ala	Leu	Leu	Ala	Gln	Glu	Gln	Val	Ser	Phe	Phe	Glu	Glu
	130					135					140				
Glu	Arg	Asn	Ile	Leu	Ser	Arg	Ser	Thr	Ser	Pro	Trp	Ile	Pro	Gln	Leu
145				150						155					160
Gln	Tyr	Ala	Phe	Gln	Asp	Lys	Asn	His	Leu	Tyr	Leu	Val	Met	Glu	Tyr
			165						170					175	
Gln	Pro	Gly	Gly	Asp	Leu	Leu	Ser	Leu	Leu	Asn	Arg	Tyr	Glu	Asp	Gln
		180						185					190		
Leu	Asp	Glu	Asn	Leu	Ile	Gln	Phe	Tyr	Leu	Ala	Glu	Leu	Ile	Leu	Ala
	195					200					205				
Val	His	Ser	Val	His	Leu	Met	Gly	Tyr	Val	His	Arg	Asp	Ile	Lys	Pro
	210					215					220				
Glu	Asn	Ile	Leu	Val	Asp	Arg	Thr	Gly	His	Ile	Lys	Leu	Val	Asp	Phe
225				230						235					240
Gly	Ser	Ala	Ala	Lys	Met	Asn	Ser	Asn	Lys	Val	Asn	Ala	Lys	Leu	Pro
			245						250					255	
Ile	Gly	Thr	Pro	Asp	Tyr	Met	Ala	Pro	Glu	Val	Leu	Thr	Val	Met	Asn

260					265					270						
Gly	Asp	Gly	Lys	Gly	Thr	Tyr	Gly	Leu	Asp	Cys	Asp	Trp	Trp	Ser	Val	
275					280					285						
Gly	Val	Ile	Ala	Tyr	Glu	Met	Ile	Tyr	Gly	Arg	Ser	Pro	Phe	Ala	Glu	
290					295					300						
Gly	Thr	Ser	Ala	Arg	Thr	Phe	Asn	Asn	Ile	Met	Asn	Phe	Gln	Arg	Phe	
305					310					315					320	
Leu	Lys	Phe	Pro	Asp	Asp	Pro	Lys	Val	Ser	Ser	Asp	Phe	Leu	Asp	Leu	
					325					330					335	
Ile	Gln	Ser	Leu	Leu	Cys	Gly	Gln	Lys	Glu	Arg	Leu	Lys	Phe	Glu	Gly	
					340					345					350	
Leu	Cys	Cys	His	Pro	Phe	Phe	Ser	Lys	Ile	Asp	Trp	Asn	Asn	Ile	Arg	
355					360					365						
Asn	Ala	Pro	Pro	Pro	Phe	Val	Pro	Thr	Leu	Lys	Ser	Asp	Asp	Asp	Thr	
370					375					380						
Ser	Asn	Phe	Asp	Glu	Pro	Glu	Lys	Asn	Ser	Trp	Val	Ser	Ser	Ser	Pro	
385					390					395					400	
Cys	Gln	Leu	Ser	Pro	Ser	Gly	Phe	Ser	Gly	Glu	Glu	Leu	Pro	Phe	Val	
					405					410					415	
Gly	Phe	Ser	Tyr	Ser	Lys	Ala	Leu	Gly	Ile	Leu	Gly	Arg	Ser	Glu	Ser	
					420					425					430	
Val	Val	Ser	Gly	Leu	Asp	Ser	Pro	Ala	Lys	Thr	Ser	Ser	Met	Glu	Lys	
435					440					445						
Lys	Leu	Leu	Ile	Lys	Ser	Lys	Glu	Leu	Gln	Asp	Ser	Gln	Asp	Lys	Cys	
450					455					460						
His	Lys	Met	Glu	Gln	Glu	Met	Thr	Arg	Leu	His	Arg	Arg	Val	Ser	Glu	
465					470					475					480	
Val	Glu	Ala	Val	Leu	Ser	Gln	Lys	Glu	Val	Glu	Leu	Lys	Ala	Ser	Glu	
					485					490					495	
Thr	Gln	Arg	Ser	Leu	Leu	Glu	Gln	Asp	Leu	Ala	Thr	Tyr	Ile	Thr	Glu	
					500					505					510	
Cys	Ser	Ser	Leu	Lys	Arg	Ser	Leu	Glu	Gln	Ala	Arg	Met	Glu	Val	Ser	
515					520					525						
Gln	Glu	Asp	Asp	Lys	Ala	Leu	Gln	Leu	Leu	His	Asp	Ile	Arg	Glu	Gln	
530					535					540						
Ser	Arg	Lys	Leu	Gln	Glu	Ile	Lys	Glu	Gln	Glu	Tyr	Gln	Ala	Gln	Val	
545					550					555					560	
Glu	Glu	Met	Arg	Leu	Met	Met	Asn	Gln	Leu	Glu	Glu	Asp	Leu	Val	Ser	

565					570					575						
Ala	Arg	Arg	Arg	Ser	Asp	Leu	Tyr	Glu	Ser	Glu	Leu	Arg	Glu	Ser	Arg	
580					585					590						
Leu	Ala	Ala	Glu	Glu	Phe	Lys	Arg	Lys	Ala	Thr	Glu	Cys	Gln	His	Lys	
595					600					605						
Leu	Leu	Lys	Ala	Lys	Asp	Gln	Gly	Lys	Pro	Glu	Val	Gly	Glu	Tyr	Ala	
610					615					620						
Lys	Leu	Glu	Lys	Ile	Asn	Ala	Glu	Gln	Gln	Leu	Lys	Ile	Gln	Glu	Leu	
625					630					635					640	
Gln	Glu	Lys	Leu	Glu	Lys	Ala	Val	Lys	Ala	Ser	Thr	Glu	Ala	Thr	Glu	
645					650					655						
Leu	Leu	Gln	Asn	Ile	Arg	Gln	Ala	Lys	Glu	Arg	Ala	Glu	Arg	Glu	Leu	
660					665					670						
Glu	Lys	Leu	Gln	Asn	Arg	Glu	Asp	Ser	Ser	Glu	Gly	Ile	Arg	Lys	Lys	
675					680					685						
Leu	Val	Glu	Ala	Glu	Glu	Arg	Arg	His	Ser	Leu	Glu	Asn	Lys	Val	Lys	
690					695					700						
Arg	Leu	Glu	Thr	Met	Glu	Arg	Arg	Glu	Asn	Arg	Leu	Lys	Asp	Asp	Ile	
705					710					715					720	
Gln	Thr	Lys	Ser	Gln	Gln	Ile	Gln	Gln	Met	Ala	Asp	Lys	Ile	Leu	Glu	
725					730					735						
Leu	Glu	Glu	Lys	His	Arg	Glu	Ala	Gln	Val	Ser	Ala	Gln	His	Leu	Glu	
740					745					750						
Val	His	Leu	Lys	Gln	Lys	Glu	Gln	His	Tyr	Glu	Glu	Lys	Ile	Lys	Val	
755					760					765						
Leu	Asp	Asn	Gln	Ile	Lys	Lys	Asp	Leu	Ala	Asp	Lys	Glu	Thr	Leu	Glu	
770					775					780						
Asn	Met	Met	Gln	Arg	His	Glu	Glu	Glu	Ala	His	Glu	Lys	Gly	Lys	Ile	
785					790					795					800	
Leu	Ser	Glu	Gln	Lys	Ala	Met	Ile	Asn	Ala	Met	Asp	Ser	Lys	Ile	Arg	
805					810					815						
Ser	Leu	Glu	Gln	Arg	Ile	Val	Glu	Leu	Ser	Glu	Ala	Asn	Lys	Leu	Ala	
820					825					830						
Ala	Asn	Ser	Ser	Leu	Phe	Thr	Gln	Arg	Asn	Met	Lys	Ala	Gln	Glu	Glu	
835					840					845						
Met	Ile	Ser	Glu	Leu	Arg	Gln	Gln	Lys	Phe	Tyr	Leu	Glu	Thr	Gln	Ala	
850					855					860						
Gly	Lys	Leu	Glu	Ala	Gln	Asn	Arg	Lys	Leu	Glu	Glu	Gln	Leu	Glu	Lys	

865	870	875	880
Ile Ser His Gln Asp 885	His Ser Asp Lys Asn Arg 890	Leu Leu Glu Leu Glu 895	
Thr Arg Leu Arg Glu Val Ser Leu Glu His Glu Glu Gln Lys Leu Glu 900	905	910	
Leu Lys Arg Gln Leu Thr Glu Leu Gln Leu Ser Leu Gln Glu Arg Glu 915	920	925	
Ser Gln Leu Thr Ala Leu Gln Ala Ala Arg Ala Ala Leu Glu Ser Gln 930	935	940	
Leu Arg Gln Ala Lys Thr Glu Leu Glu Glu Thr Thr Ala Glu Ala Glu 945	950	955	960
Glu Glu Ile Gln Ala Leu Thr Ala His Arg Asp Glu Ile Gln Arg Lys 965	970	975	
Phe Asp Ala Leu Arg Asn Ser Cys Thr Val Ile Thr Asp Leu Glu Glu 980	985	990	
Gln Leu Asn Gln Leu Thr Glu Asp Asn Ala Glu Leu Asn Asn Gln Asn 995	1000	1005	
Phe Tyr Leu Ser Lys Gln Leu Asp Glu Ala Ser Gly Ala Asn Asp Glu 1010	1015	1020	
Ile Val Gln Leu Arg Ser Glu Val Asp His Leu Arg Arg Glu Ile Thr 1025	1030	1035	1040
Glu Arg Glu Met Gln Leu Thr Ser Gln Lys Gln Thr Met Glu Ala Leu 1045	1050	1055	
Lys Thr Thr Cys Thr Met Leu Glu Glu Gln Val Met Asp Leu Glu Ala 1060	1065	1070	
Leu Asn Asp Glu Leu Leu Glu Lys Glu Arg Gln Trp Glu Ala Trp Arg 1075	1080	1085	
Ser Val Leu Gly Asp Glu Lys Ser Gln Phe Glu Cys Arg Val Arg Glu 1090	1095	1100	
Leu Gln Arg Met Leu Asp Thr Glu Lys Gln Ser Arg Ala Arg Ala Asp 1105	1110	1115	1120
Gln Arg Ile Thr Glu Ser Arg Gln Val Val Glu Leu Ala Val Lys Glu 1125	1130	1135	
His Lys Ala Glu Ile Leu Ala Leu Gln Gln Ala Leu Lys Glu Gln Lys 1140	1145	1150	
Leu Lys Ala Glu Ser Leu Ser Asp Lys Leu Asn Asp Leu Glu Lys Lys 1155	1160	1165	
His Ala Met Leu Glu Met Asn Ala Arg Ser Leu Gln Gln Lys Leu Glu			

1170	1175	1180
Thr Glu Arg Glu Leu Lys Gln Arg Leu Leu Glu Glu Gln Ala Lys Leu 1185 1190 1195 1200		
Gln Gln Gln Met Asp Leu Gln Lys Asn His Ile Phe Arg Leu Thr Gln 1205 1210 1215		
Gly Leu Gln Glu Ala Leu Asp Arg Ala Asp Leu Leu Lys Thr Glu Arg 1220 1225 1230		
Ser Asp Leu Glu Tyr Gln Leu Glu Asn Ile Gln Val Leu Tyr Ser His 1235 1240 1245		
Glu Lys Val Lys Met Glu Gly Thr Ile Ser Gln Gln Thr Lys Leu Ile 1250 1255 1260		
Asp Phe Leu Gln Ala Lys Met Asp Gln Pro Ala Lys Lys Lys Lys Val 1265 1270 1275 1280		
Pro Leu Gln Tyr Asn Glu Leu Lys Leu Ala Leu Glu Lys Glu Lys Ala 1285 1290 1295		
Arg Cys Ala Glu Leu Glu Glu Ala Leu Gln Lys Thr Arg Ile Glu Leu 1300 1305 1310		
Arg Ser Ala Arg Glu Glu Ala Ala His Arg Lys Ala Thr Asp His Pro 1315 1320 1325		
His Pro Ser Thr Pro Ala Thr Ala Arg Gln Gln Ile Ala Met Ser Ala 1330 1335 1340		
Ile Val Arg Ser Pro Glu His Gln Pro Ser Ala Met Ser Leu Leu Ala 1345 1350 1355 1360		
Pro Pro Ser Ser Arg Arg Lys Glu Ser Ser Thr Pro Glu Glu Phe Ser 1365 1370 1375		
Arg Arg Leu Lys Glu Arg Met His His Asn Ile Pro His Arg Phe Asn 1380 1385 1390		
Val Gly Leu Asn Met Arg Ala Thr Lys Cys Ala Val Cys Leu Asp Thr 1395 1400 1405		
Val His Phe Gly Arg Gln Ala Ser Lys Cys Leu Glu Cys Gln Val Met 1410 1415 1420		
Cys His Pro Lys Cys Ser Thr Cys Leu Pro Ala Thr Cys Gly Leu Pro 1425 1430 1435 1440		
Ala Glu Tyr Ala Thr His Phe Thr Glu Ala Phe Cys Arg Asp Lys Met 1445 1450 1455		
Asn Ser Pro Gly Leu Gln Thr Lys Glu Pro Ser Ser Ser Leu His Leu 1460 1465 1470		
Glu Gly Trp Met Lys Val Pro Arg Asn Asn Lys Arg Gly Gln Gln Gly		

1475	1480	1485
Trp Asp Arg Lys Tyr Ile Val Leu Glu Gly Ser Lys Val Leu Ile Tyr 1490	1495	1500
Asp Asn Glu Ala Arg Glu Ala Gly Gln Arg Pro Val Glu Glu Phe Glu 1505	1510	1515 1520
Leu Cys Leu Pro Asp Gly Asp Val Ser Ile His Gly Ala Val Gly Ala 1525	1530	1535
Ser Glu Leu Ala Asn Thr Ala Lys Ala Asp Val Pro Tyr Ile Leu Lys 1540	1545	1550
Met Glu Ser His Pro His Thr Thr Cys Trp Pro Gly Arg Thr Leu Tyr 1555	1560	1565
Leu Leu Ala Pro Ser Phe Pro Asp Lys Gln Arg Trp Val Thr Ala Leu 1570	1575	1580
Glu Ser Val Val Ala Gly Gly Arg Val Ser Arg Glu Lys Ala Glu Ala 1585	1590	1595 1600
Asp Ala Lys Leu Leu Gly Asn Ser Leu Leu Lys Leu Glu Gly Asp Asp 1605	1610	1615
Arg Leu Asp Met Asn Cys Thr Leu Pro Phe Ser Asp Gln Val Val Leu 1620	1625	1630
Val Gly Thr Glu Glu Gly Leu Tyr Ala Leu Asn Val Leu Lys Asn Ser 1635	1640	1645
Leu Thr His Val Pro Gly Ile Gly Ala Val Phe Gln Ile Tyr Ile Ile 1650	1655	1660
Lys Asp Leu Glu Lys Leu Leu Met Ile Ala Gly Glu Glu Arg Ala Leu 1665	1670	1675 1680
Cys Leu Val Asp Val Lys Lys Val Lys Gln Ser Leu Ala Gln Ser His 1685	1690	1695
Leu Pro Ala Gln Pro Asp Ile Ser Pro Asn Ile Phe Glu Ala Val Lys 1700	1705	1710
Gly Cys His Leu Phe Gly Ala Gly Lys Ile Glu Asn Gly Leu Cys Ile 1715	1720	1725
Cys Ala Ala Met Pro Ser Lys Val Val Ile Leu Arg Tyr Asn Glu Asn 1730	1735	1740
Leu Ser Lys Tyr Cys Ile Arg Lys Glu Ile Glu Thr Ser Glu Pro Cys 1745	1750	1755 1760
Ser Cys Ile His Phe Thr Asn Tyr Ser Ile Leu Ile Gly Thr Asn Lys 1765	1770	1775
Phe Tyr Glu Ile Asp Met Lys Gln Tyr Thr Leu Glu Glu Phe Leu Asp		

1780	1785	1790
Lys Asn Asp His Ser Leu Ala Pro Ala Val Phe Ala Ala Ser Ser Asn 1795 1800 1805		
Ser Phe Pro Val Ser Ile Val Gln Val Asn Ser Ala Gly Gln Arg Glu 1810 1815 1820		
Glu Tyr Leu Leu Cys Phe His Glu Phe Gly Val Phe Val Asp Ser Tyr 1825 1830 1835 1840		
Gly Arg Arg Ser Arg Thr Asp Asp Leu Lys Trp Ser Arg Leu Pro Leu 1845 1850 1855		
Ala Phe Ala Tyr Arg Glu Pro Tyr Leu Phe Val Thr His Phe Asn Ser 1860 1865 1870		
Leu Glu Val Ile Glu Ile Gln Ala Arg Ser Ser Ala Gly Thr Pro Ala 1875 1880 1885		
Arg Ala Tyr Leu Asp Ile Pro Asn Pro Arg Tyr Leu Gly Pro Ala Ile 1890 1895 1900		
Ser Ser Gly Ala Ile Tyr Leu Ala Ser Ser Tyr Gln Asp Lys Leu Arg 1905 1910 1915 1920		
Val Ile Cys Cys Lys Gly Asn Leu Val Lys Glu Ser Gly Thr Glu His 1925 1930 1935		
His Arg Gly Pro Ser Thr Ser Arg Ser Ser Pro Asn Lys Arg Gly Pro 1940 1945 1950		
Pro Thr Tyr Asn Glu His Ile Thr Lys Arg Val Ala Ser Ser Pro Ala 1955 1960 1965		
Pro Pro Glu Gly Pro Ser His Pro Arg Glu Pro Ser Thr Pro His Arg 1970 1975 1980		
Tyr Arg Glu Gly Arg Thr Glu Leu Arg Arg Asp Lys Ser Pro Gly Arg 1985 1990 1995 2000		
Pro Leu Glu Arg Glu Lys Ser Pro Gly Arg Met Leu Ser Thr Arg Arg 2005 2010 2015		
Glu Arg Ser Pro Gly Arg Leu Phe Glu Asp Ser Ser Arg Gly Arg Leu 2020 2025 2030		
Pro Ala Gly Ala Val Arg Thr Pro Leu Ser Gln Val Asn Lys Val Trp 2035 2040 2045		
Asp Gln Ser Ser Val 2050		

<210> 12
 <211> 5691
 <212> DNA

<213> Homo sapiens

<400> 12

```
atgaaagcca tgccttgaa ctggacctgc cttctctccc acctcctcat ggtgggcatg 60
ggctcctcca ctttgcacac ccggcagcca gccccgtgt cccagaagca gcggtcattt 120
gtcacattcc gaggagagcc cggcaggggt ttcaatcacc tgggtgggga tgagaggaca 180
ggacacattt acttgggggc cgtcaatcgg atttacaagc tctccagega cctgaaggtc 240
ttggtgacgc atgagacagg gccggacgag gacaaccca agtggttacc accccgcac 300
gtccagacct gcaatgagcc cctgaccacc accaacaatg tcaacaagat gctcctcata 360
gactacaagg agaacaggct gattgcctgt gggagcctgt accaaggcat ctgcaagctg 420
ctgaggctgg aggacctctt caagctgggg gagccttacc ataagaagga gcactatctg 480
tcaggtgtca acgagagcgg ctgagctctt ggagtgatcg tctcctacag caacctggat 540
gacaagctgt tcattgccac ggcagtggtt ggggaagccc agtattttcc caccatctcc 600
agccggaaac tgaccaagaa ctctgaggcg gatggcatgt tcgcgtacgt cttccatgat 660
gagttcgtgg cctcgatgat taagatccct tcggacacct tcaccatcat ccctgacttt 720
gatatctact atgtctatgg ttttagcagt ggcaactttg tctacttttt gaccctccaa 780
cctgagatgg tgtctcacc aggtccacc accaaggagc aggtgtatac atccaagctc 840
gtgaggcttt gcaaggagga cacagccttc aactcctatg tagaggtgcc cattggctgt 900
gagcgcagtg ggggtggagta ccgcctgctg caggctgcct acctgtccaa agcggggggc 960
gtgcttgcca ggaccttg agtccatcca gatgatgacc tgctcttcac cgtcttctcc 1020
aagggccaga agcggaaaat gaaatccctg gatgagtcgg ccctgtgcat cttcatcttg 1080
aagcagataa atgaccgcat taaggagcgg ctgcagtctt gttaccgggg cgagggcacg 1140
ctggacctgg cctggctcaa ggtgaaggac atccccctgca gcagtgcgct ctttaaccatt 1200
gacgataact tctgtggcct ggacatgaat gctccccctg gagtgtccga catggtgctg 1260
ggaattcccc tcttcacgga ggacagggac cgcagtacgt ctgtcatcgc atatgtctac 1320
aagaaccact ctctggcctt tgtgggcacc aaaagtggca agctgaagaa gatccgggtg 1380
gatggacca ggggcaacgc cctccagtat gagacgggtg aggtggtgga ccccgggcca 1440
gtcctccggg atatggcctt ctccaaggac cacgagcaac tctacatcat gtcagagagg 1500
cagctacca gagtccctgt ggagtcctgt ggtcagtatc agagctgcgg cgagtgcctt 1560
ggctcaggcg acccccactg tggctgggtg gtgctgcaca acacgtgcac ccggaaggag 1620
cgggtgtgagc ggtccaagga gccccgcagg tttgcctcgg agatgaagca gtgtgtccgg 1680
ctgacggtcc atcccaacaa tatctccgtc tctcagtaca acctgtgctg ggtcctggag 1740
acgtacaatg tcccggagct gtcagctggc gtcaactgca ccttgaggga cctgtcagag 1800
atggatgggc tggctcgtgg caatcagatc cagtgcctact cccctgcagc caaggagggtg 1860
ccccgatca tcacagagaa tggggaccac catgtcgtac agcttcagct caaatcaaag 1920
gagaccggca tgaccttcgc cagcaccagc tttgtcttct acaattgcag cgtccacaat 1980
tcgtgcctgt cctgcgtgga gagtccatac cgctgccact ggtgtaaata ccggcatgtc 2040
tgacccatg accccaagac ctgctccttc caggaaggcc gagtgaagct gcccgaggac 2100
tgccccagc tgctgcgagt ggacaagatc ctggtgcccg tggagggtgat caagcctatc 2160
acgtgaagg ccaagaacct cccccagccc cagtctgggc agcgtggcta cgaatgcac 2220
ctcaacattc agggcagcga gcagcgagt cccgcctgc gcttcaacag ctccagcgta 2280
cagtgcaga acacctctta ttcctatgaa gggatggaga tcaacaacct gccctggag 2340
ttgacagtcg tgtggaatgg gcacttcaac attgacaacc cagctcagaa taaagtccac 2400
ctctacaagt gtggagccat gcgtgagagc tgcgggctgt gcctcaaggc tgaccagac 2460
ttcgcatgtg gctggtgcca gggcccaggc cagtgcacc tgcgccagca ctgccctgcc 2520
caggagagcc agtggctgga gctgtctggt gccaaaagca agtgcacaaa cccccgcac 2580
acagagataa tcccgtgac agggccccgg gaagggggca ccaaggctac tatccgagg 2640
gagaacctgg gcctggaatt tcgcgacatc gcctcccatg tcaagggtgc tggcggtgag 2700
tgacgccctt tagtgatgg ttacatccct gcagaacaga tcgtgtgtga gatgggggag 2760
gccaaagcca gccagcatgc aggtctcgtg gagatctgcg tggctgtgtg tcggcctgaa 2820
ttcatggccc ggtcctcaca gctctattac ttcatgacac tgactctctc agatctgaag 2880
cccagccggg gggccatgtc cggaggggac caagtgaaca tcacaggcac caacctgaat 2940
gccggaagca acgtgggtgg gatgtttgga aagcagccct gtctcttcca caggcgatct 3000
ccatcctaca ttgtctgcaa caccacatcc tcagatgagg tgctagagat gaagggtgcg 3060
gtgcaggtgg acagggccaa gatccaccag gacctggtct ttcagtatgt ggaagacccc 3120
accatcgtgc ggattgagcc agaatggagc attgtcagtg gaaacacacc catcgccgta 3180
tgggggaccc acctggacct catacagaac cccagatcc gtgccaagca tggagggaag 3240
```

```

gagcacatca atatctgtga ggttctgaac gctactgaga tgacctgtca ggcgcccgcc 3300
ctcgcctctgg gtcctgacca ccagtcagac ctgaccgaga ggcccagga gtttggttc 3360
atcctggaca acgtccagtc cctgctcatc ctcaacaaga ccaacttcac ctactatccc 3420
aaccgggtgt ttgaggcctt tggtcctca ggaatcctgg agctcaagcc tggcacgccc 3480
atcatcctaa agggcaagaa cctgatcccg cctgtggctg ggggcaacgt gaagctgaac 3540
tacactgtgc tgggtgggga gaagccgtgc accgtgaccg tgtcagatgt ccagctgctc 3600
tgcgagtccc ccaacctcat cggcaggcac aaagtgatgg cccgtgtcgg tggcatggag 3660
tactccccgg ggatggtgta cattgccccg gacagcccgc tcagcctgcc cgccatcgtc 3720
agcatcgagc tggttgggcg cctcctcatc attttcatcg tggccgtgct cattgcctat 3780
aaacgcaagt cccgcgaaag tgacctcacg ctgaagcggc tgcagatgca gatggacaac 3840
ctggagtccc gtgtggccct ggagtgcagc gaagcctttg ccgagctgca gacggacatc 3900
catgagctga ccagtgcact ggatggagcc gggattccgt tcctggacta tagaacttac 3960
accatgctgg tgcgtgtccc aggaattgaa gaccacctg tcctccggga ccttgaggtc 4020
ccgggctacc ggcaggagcg tgtggagaaa ggctgaagc tcttcgccc gctcatcaac 4080
aacaaggtgt tcctgctgtc cttcatccgc acgcttgagt cccagcgtag cttctccatg 4140
cgcgaccgtg gcaacgtggc ctactcatc atgaccgtgc tgcagagcaa gctggagtac 4200
gccactgatg tgctgaagca gctgctggcc gacctcattg acaagaacct ggagagcaag 4260
aaccacccta agctgctgct caggaggact gagtcatggg ctgagaagat gctgaccaat 4320
tggtttactt tcctcctcta caagttcctc aaggagtgtg ctggggagcc cctcttctcc 4380
ctgttctgtg ccatcaagca gcagatggag aagggcccca ttgacgccat cacgggagag 4440
gcccgtactc ccttgagcga ggacaagctc atccgccagc agattgacta caaaaccttg 4500
gtcctgagct gtgtcagccc agacaatgcc aacagccccg aggtcccagt aaagatcctc 4560
aactgtgaca ccatcactca ggtcaaggag aagattctgg atgccatctt caagaatgtg 4620
ccttgctccc accggcccaa agctgcagat atggatctgg agtggcgaca aggaagtggg 4680
gcaaggatga tcttgagga tgaagacatc accaccaaga ttgagaatga ttggaagcga 4740
ctgaacacac tggccacta ccaggtgcca gatggttccg tgggtggcatt agtgtccaag 4800
caggtgacag cctataacgc agtgaacaac tccaccgtct ccaggacctc agcaagtaaa 4860
tatgaaaaca tgatccggta cacgggcagc cccgacagcc tccgctcacg gacacctatg 4920
atcactcctg acctggagag tggagtcaag atgtggcacc tagtgaagaa ccacgagcac 4980
ggagaccaga aggaggggga ccgggggagc aagatggtgt ctgaaatcta cctgacctga 5040
ctcctggcca ctaagggcac actgcagaag tttgtggatg acctctttga gaccatcttc 5100
agcacggcac accgtggctc tgccctgccc ctggccatca agtacatgtt tgacttctctg 5160
gatgagcagg ctgataaaca tggcattcat gacccgcacg tccgccatac ctggaagagc 5220
aattgcctgc cctgaggtt ttgggtcaac atgatcaaga acccgagtt tgtgtttgac 5280
atccataaga acagcatcac agacgcctgc ctctctgtgg tggctcagac cttcatggac 5340
tcttgctcca cgtcagagca cgggctgggc aaggactcgc cctccaacaa gctgctgtat 5400
gccaaggaca tccccagcta caagaattgg gtggagaggt attactcaga catagggaag 5460
atgccagcca tcagcgacca agaactgaac gcatacctgg ctgagcagtc ccggatgcac 5520
atgaatgagt tcaacaccat gagtgcactc tcagagatct tctcctatgt gggcaaatac 5580
agcgaggaga tccttgagcc tctggaccac gatgaccagt gtgggaagca gaaactggcc 5640
tacaactag aacaagtcac aacctcatg agcttagaca gctgaaataa a 5691

```

<210> 13

<211> 1896

<212> PRT

<213> Homo sapiens

<400> 13

```

Met Lys Ala Met Pro Trp Asn Trp Thr Cys Leu Leu Ser His Leu Leu
  1                      5                      10                      15

```

```

Met Val Gly Met Gly Ser Ser Thr Leu Leu Thr Arg Gln Pro Ala Pro
  20                      25                      30

```

```

Leu Ser Gln Lys Gln Arg Ser Phe Val Thr Phe Arg Gly Glu Pro Ala
  35                      40                      45

```

Glu Gly Phe Asn His Leu Val Val Asp Glu Arg Thr Gly His Ile Tyr
 50 55 60
 Leu Gly Ala Val Asn Arg Ile Tyr Lys Leu Ser Ser Asp Leu Lys Val
 65 70 75 80
 Leu Val Thr His Glu Thr Gly Pro Asp Glu Asp Asn Pro Lys Cys Tyr
 85 90 95
 Pro Pro Arg Ile Val Gln Thr Cys Asn Glu Pro Leu Thr Thr Thr Asn
 100 105 110
 Asn Val Asn Lys Met Leu Leu Ile Asp Tyr Lys Glu Asn Arg Leu Ile
 115 120 125
 Ala Cys Gly Ser Leu Tyr Gln Gly Ile Cys Lys Leu Leu Arg Leu Glu
 130 135 140
 Asp Leu Phe Lys Leu Gly Glu Pro Tyr His Lys Lys Glu His Tyr Leu
 145 150 155 160
 Ser Gly Val Asn Glu Ser Gly Ser Val Phe Gly Val Ile Val Ser Tyr
 165 170 175
 Ser Asn Leu Asp Asp Lys Leu Phe Ile Ala Thr Ala Val Asp Gly Lys
 180 185 190
 Pro Glu Tyr Phe Pro Thr Ile Ser Ser Arg Lys Leu Thr Lys Asn Ser
 195 200 205
 Glu Ala Asp Gly Met Phe Ala Tyr Val Phe His Asp Glu Phe Val Ala
 210 215 220
 Ser Met Ile Lys Ile Pro Ser Asp Thr Phe Thr Ile Ile Pro Asp Phe
 225 230 235 240
 Asp Ile Tyr Tyr Val Tyr Gly Phe Ser Ser Gly Asn Phe Val Tyr Phe
 245 250 255
 Leu Thr Leu Gln Pro Glu Met Val Ser Pro Pro Gly Ser Thr Thr Lys
 260 265 270
 Glu Gln Val Tyr Thr Ser Lys Leu Val Arg Leu Cys Lys Glu Asp Thr
 275 280 285
 Ala Phe Asn Ser Tyr Val Glu Val Pro Ile Gly Cys Glu Arg Ser Gly
 290 295 300
 Val Glu Tyr Arg Leu Leu Gln Ala Ala Tyr Leu Ser Lys Ala Gly Ala
 305 310 315 320
 Val Leu Gly Arg Thr Leu Gly Val His Pro Asp Asp Asp Leu Leu Phe
 325 330 335
 Thr Val Phe Ser Lys Gly Gln Lys Arg Lys Met Lys Ser Leu Asp Glu
 340 345 350

Ser Ala Leu Cys Ile Phe Ile Leu Lys Gln Ile Asn Asp Arg Ile Lys
 355 360 365
 Glu Arg Leu Gln Ser Cys Tyr Arg Gly Glu Gly Thr Leu Asp Leu Ala
 370 375 380
 Trp Leu Lys Val Lys Asp Ile Pro Cys Ser Ser Ala Leu Leu Thr Ile
 385 390 395 400
 Asp Asp Asn Phe Cys Gly Leu Asp Met Asn Ala Pro Leu Gly Val Ser
 405 410 415
 Asp Met Val Arg Gly Ile Pro Val Phe Thr Glu Asp Arg Asp Arg Met
 420 425 430
 Thr Ser Val Ile Ala Tyr Val Tyr Lys Asn His Ser Leu Ala Phe Val
 435 440 445
 Gly Thr Lys Ser Gly Lys Leu Lys Lys Ile Arg Val Asp Gly Pro Arg
 450 455 460
 Gly Asn Ala Leu Gln Tyr Glu Thr Val Gln Val Val Asp Pro Gly Pro
 465 470 475 480
 Val Leu Arg Asp Met Ala Phe Ser Lys Asp His Glu Gln Leu Tyr Ile
 485 490 495
 Met Ser Glu Arg Gln Leu Thr Arg Val Pro Val Glu Ser Cys Gly Gln
 500 505 510
 Tyr Gln Ser Cys Gly Glu Cys Leu Gly Ser Gly Asp Pro His Cys Gly
 515 520 525
 Trp Cys Val Leu His Asn Thr Cys Thr Arg Lys Glu Arg Cys Glu Arg
 530 535 540
 Ser Lys Glu Pro Arg Arg Phe Ala Ser Glu Met Lys Gln Cys Val Arg
 545 550 555 560
 Leu Thr Val His Pro Asn Asn Ile Ser Val Ser Gln Tyr Asn Val Leu
 565 570 575
 Leu Val Leu Glu Thr Tyr Asn Val Pro Glu Leu Ser Ala Gly Val Asn
 580 585 590
 Cys Thr Phe Glu Asp Leu Ser Glu Met Asp Gly Leu Val Val Gly Asn
 595 600 605
 Gln Ile Gln Cys Tyr Ser Pro Ala Ala Lys Glu Val Pro Arg Ile Ile
 610 615 620
 Thr Glu Asn Gly Asp His His Val Val Gln Leu Gln Leu Lys Ser Lys
 625 630 635 640
 Glu Thr Gly Met Thr Phe Ala Ser Thr Ser Phe Val Phe Tyr Asn Cys
 645 650 655

Ser Val His Asn Ser Cys Leu Ser Cys Val Glu Ser Pro Tyr Arg Cys
 660 665 670
 His Trp Cys Lys Tyr Arg His Val Cys Thr His Asp Pro Lys Thr Cys
 675 680 685
 Ser Phe Gln Glu Gly Arg Val Lys Leu Pro Glu Asp Cys Pro Gln Leu
 690 695 700
 Leu Arg Val Asp Lys Ile Leu Val Pro Val Glu Val Ile Lys Pro Ile
 705 710 715 720
 Thr Leu Lys Ala Lys Asn Leu Pro Gln Pro Gln Ser Gly Gln Arg Gly
 725 730 735
 Tyr Glu Cys Ile Leu Asn Ile Gln Gly Ser Glu Gln Arg Val Pro Ala
 740 745 750
 Leu Arg Phe Asn Ser Ser Ser Val Gln Cys Gln Asn Thr Ser Tyr Ser
 755 760 765
 Tyr Glu Gly Met Glu Ile Asn Asn Leu Pro Val Glu Leu Thr Val Val
 770 775 780
 Trp Asn Gly His Phe Asn Ile Asp Asn Pro Ala Gln Asn Lys Val His
 785 790 795 800
 Leu Tyr Lys Cys Gly Ala Met Arg Glu Ser Cys Gly Leu Cys Leu Lys
 805 810 815
 Ala Asp Pro Asp Phe Ala Cys Gly Trp Cys Gln Gly Pro Gly Gln Cys
 820 825 830
 Thr Leu Arg Gln His Cys Pro Ala Gln Glu Ser Gln Trp Leu Glu Leu
 835 840 845
 Ser Gly Ala Lys Ser Lys Cys Thr Asn Pro Arg Ile Thr Glu Ile Ile
 850 855 860
 Pro Val Thr Gly Pro Arg Glu Gly Gly Thr Lys Val Thr Ile Arg Gly
 865 870 875 880
 Glu Asn Leu Gly Leu Glu Phe Arg Asp Ile Ala Ser His Val Lys Val
 885 890 895
 Ala Gly Val Glu Cys Ser Pro Leu Val Asp Gly Tyr Ile Pro Ala Glu
 900 905 910
 Gln Ile Val Cys Glu Met Gly Glu Ala Lys Pro Ser Gln His Ala Gly
 915 920 925
 Phe Val Glu Ile Cys Val Ala Val Cys Arg Pro Glu Phe Met Ala Arg
 930 935 940
 Ser Ser Gln Leu Tyr Tyr Phe Met Thr Leu Thr Leu Ser Asp Leu Lys
 945 950 955 960

Pro Ser Arg Gly Pro Met Ser Gly Gly Thr Gln Val Thr Ile Thr Gly
 965 970 975
 Thr Asn Leu Asn Ala Gly Ser Asn Val Val Val Met Phe Gly Lys Gln
 980 985 990
 Pro Cys Leu Phe His Arg Arg Ser Pro Ser Tyr Ile Val Cys Asn Thr
 995 1000 1005
 Thr Ser Ser Asp Glu Val Leu Glu Met Lys Val Ser Val Gln Val Asp
 1010 1015 1020
 Arg Ala Lys Ile His Gln Asp Leu Val Phe Gln Tyr Val Glu Asp Pro
 1025 1030 1035 1040
 Thr Ile Val Arg Ile Glu Pro Glu Trp Ser Ile Val Ser Gly Asn Thr
 1045 1050 1055
 Pro Ile Ala Val Trp Gly Thr His Leu Asp Leu Ile Gln Asn Pro Gln
 1060 1065 1070
 Ile Arg Ala Lys His Gly Gly Lys Glu His Ile Asn Ile Cys Glu Val
 1075 1080 1085
 Leu Asn Ala Thr Glu Met Thr Cys Gln Ala Pro Ala Leu Ala Leu Gly
 1090 1095 1100
 Pro Asp His Gln Ser Asp Leu Thr Glu Arg Pro Glu Glu Phe Gly Phe
 1105 1110 1115 1120
 Ile Leu Asp Asn Val Gln Ser Leu Leu Ile Leu Asn Lys Thr Asn Phe
 1125 1130 1135
 Thr Tyr Tyr Pro Asn Pro Val Phe Glu Ala Phe Gly Pro Ser Gly Ile
 1140 1145 1150
 Leu Glu Leu Lys Pro Gly Thr Pro Ile Ile Leu Lys Gly Lys Asn Leu
 1155 1160 1165
 Ile Pro Pro Val Ala Gly Gly Asn Val Lys Leu Asn Tyr Thr Val Leu
 1170 1175 1180
 Val Gly Glu Lys Pro Cys Thr Val Thr Val Ser Asp Val Gln Leu Leu
 1185 1190 1195 1200
 Cys Glu Ser Pro Asn Leu Ile Gly Arg His Lys Val Met Ala Arg Val
 1205 1210 1215
 Gly Gly Met Glu Tyr Ser Pro Gly Met Val Tyr Ile Ala Pro Asp Ser
 1220 1225 1230
 Pro Leu Ser Leu Pro Ala Ile Val Ser Ile Ala Val Ala Gly Gly Leu
 1235 1240 1245
 Leu Ile Ile Phe Ile Val Ala Val Leu Ile Ala Tyr Lys Arg Lys Ser
 1250 1255 1260

Arg Glu Ser Asp Leu Thr Leu Lys Arg Leu Gln Met Gln Met Asp Asn
 1265 1270 1275 1280

Leu Glu Ser Arg Val Ala Leu Glu Cys Lys Glu Ala Phe Ala Glu Leu
 1285 1290 1295

Gln Thr Asp Ile His Glu Leu Thr Ser Asp Leu Asp Gly Ala Gly Ile
 1300 1305 1310

Pro Phe Leu Asp Tyr Arg Thr Tyr Thr Met Arg Val Leu Phe Pro Gly
 1315 1320 1325

Ile Glu Asp His Pro Val Leu Arg Asp Leu Glu Val Pro Gly Tyr Arg
 1330 1335 1340

Gln Glu Arg Val Glu Lys Gly Leu Lys Leu Phe Ala Gln Leu Ile Asn
 1345 1350 1355 1360

Asn Lys Val Phe Leu Leu Ser Phe Ile Arg Thr Leu Glu Ser Gln Arg
 1365 1370 1375

Ser Phe Ser Met Arg Asp Arg Gly Asn Val Ala Ser Leu Ile Met Thr
 1380 1385 1390

Val Leu Gln Ser Lys Leu Glu Tyr Ala Thr Asp Val Leu Lys Gln Leu
 1395 1400 1405

Leu Ala Asp Leu Ile Asp Lys Asn Leu Glu Ser Lys Asn His Pro Lys
 1410 1415 1420

Leu Leu Leu Arg Arg Thr Glu Ser Val Ala Glu Lys Met Leu Thr Asn
 1425 1430 1435 1440

Trp Phe Thr Phe Leu Leu Tyr Lys Phe Leu Lys Glu Cys Ala Gly Glu
 1445 1450 1455

Pro Leu Phe Ser Leu Phe Cys Ala Ile Lys Gln Gln Met Glu Lys Gly
 1460 1465 1470

Pro Ile Asp Ala Ile Thr Gly Glu Ala Arg Tyr Ser Leu Ser Glu Asp
 1475 1480 1485

Lys Leu Ile Arg Gln Gln Ile Asp Tyr Lys Thr Leu Val Leu Ser Cys
 1490 1495 1500

Val Ser Pro Asp Asn Ala Asn Ser Pro Glu Val Pro Val Lys Ile Leu
 1505 1510 1515 1520

Asn Cys Asp Thr Ile Thr Gln Val Lys Glu Lys Ile Leu Asp Ala Ile
 1525 1530 1535

Phe Lys Asn Val Pro Cys Ser His Arg Pro Lys Ala Ala Asp Met Asp
 1540 1545 1550

Leu Glu Trp Arg Gln Gly Ser Gly Ala Arg Met Ile Leu Gln Asp Glu
 1555 1560 1565

Asp Ile Thr Thr Lys Ile Glu Asn Asp Trp Lys Arg Leu Asn Thr Leu
 1570 1575 1580

Ala His Tyr Gln Val Pro Asp Gly Ser Val Val Ala Leu Val Ser Lys
 1585 1590 1595 1600

Gln Val Thr Ala Tyr Asn Ala Val Asn Asn Ser Thr Val Ser Arg Thr
 1605 1610 1615

Ser Ala Ser Lys Tyr Glu Asn Met Ile Arg Tyr Thr Gly Ser Pro Asp
 1620 1625 1630

Ser Leu Arg Ser Arg Thr Pro Met Ile Thr Pro Asp Leu Glu Ser Gly
 1635 1640 1645

Val Lys Met Trp His Leu Val Lys Asn His Glu His Gly Asp Gln Lys
 1650 1655 1660

Glu Gly Asp Arg Gly Ser Lys Met Val Ser Glu Ile Tyr Leu Thr Arg
 1665 1670 1675 1680

Leu Leu Ala Thr Lys Gly Thr Leu Gln Lys Phe Val Asp Asp Leu Phe
 1685 1690 1695

Glu Thr Ile Phe Ser Thr Ala His Arg Gly Ser Ala Leu Pro Leu Ala
 1700 1705 1710

Ile Lys Tyr Met Phe Asp Phe Leu Asp Glu Gln Ala Asp Lys His Gly
 1715 1720 1725

Ile His Asp Pro His Val Arg His Thr Trp Lys Ser Asn Cys Leu Pro
 1730 1735 1740

Leu Arg Phe Trp Val Asn Met Ile Lys Asn Pro Gln Phe Val Phe Asp
 1745 1750 1755 1760

Ile His Lys Asn Ser Ile Thr Asp Ala Cys Leu Ser Val Val Ala Gln
 1765 1770 1775

Thr Phe Met Asp Ser Cys Ser Thr Ser Glu His Arg Leu Gly Lys Asp
 1780 1785 1790

Ser Pro Ser Asn Lys Leu Leu Tyr Ala Lys Asp Ile Pro Ser Tyr Lys
 1795 1800 1805

Asn Trp Val Glu Arg Tyr Tyr Ser Asp Ile Gly Lys Met Pro Ala Ile
 1810 1815 1820

Ser Asp Gln Asp Met Asn Ala Tyr Leu Ala Glu Gln Ser Arg Met His
 1825 1830 1835 1840

Met Asn Glu Phe Asn Thr Met Ser Ala Leu Ser Glu Ile Phe Ser Tyr
 1845 1850 1855

Val Gly Lys Tyr Ser Glu Glu Ile Leu Gly Pro Leu Asp His Asp Asp
 1860 1865 1870

Gln Cys Gly Lys Gln Lys Leu Ala Tyr Lys Leu Glu Gln Val Ile Thr
 1875 1880 1885

Leu Met Ser Leu Asp Ser Asn Lys
 1890 1895

<210> 14
 <211> 1535
 <212> DNA
 <213> Homo sapiens

<400> 14
 cccgaaatgc tgccgccaag gagcaacgac accgcgtacc cggggcagtt agcgctatac 60
 cagcagctgg cgaggggaa tgccgtgggg ggctcggcgg gggcaccgcc actggggccc 120
 gtgcaggtgg tcaccgcctg cctgctgacc ctactcgtca tctggacctt gctgggcaac 180
 gtgctggtgt ccgcagccat cgtgtggagc cgccacctgc gcgccaagat gaccaacgtc 240
 ttcacgtgt ctctacctgt gtcagacctc ttcgtggcgc tgctggtcat gtcctggaag 300
 gcagtcgccc aggtggccgg ttactggccc tttgaagcgt tctgcgacgt ctgggtggcc 360
 ttcgacatca tgtgctccac cgctccatc ctgaacctgt gcgtcatcag cgtggcccgc 420
 tactgggcca tctccaggcc cttccgctac gagcgcaaga tgaccagcgc catggccttg 480
 gtcattggtcc gcccggcctg gaccttgacc agcctcatct ccttcattcc ggtccagctc 540
 aactggcaca gggaccaggc ggtctcttgg ggtgggctgg acctgccaaa caacctggcc 600
 aactggacgc cctgggagga ggccgtttgg gagcccgacg tgagggcaga gaactgtgac 660
 tccagcctga atcgaaacct cgccatccct tcctcgctca tcagcttcta catccccatg 720
 gccatcatga tcgtgacctc cagcgcctac taccgcctgc cccaggtgca gatccgcagg 780
 atttcctccc tggagagggc cgcagagcac gtgcagagct gccggagcag cgcaggctgc 840
 acgcccagca ccagcctgcg gttttccatc aagaaggaga ccgaggttct caagaccctg 900
 tcggtgatca tgggggtctt cgtgtgttgc tggctgccct tcttcacatc taactgcatg 960
 gttcctttct gcagtggaca ccccaaaggc cctccggccg gcttccccctg cgtcagttag 1020
 accacattcg atgtcttcat ctggttctgc tgggccaact cctcactcaa cccagtcccc 1080
 agtcactatg ccttcaacgc cgacttccgg aaggtgtttg cccagctgct ggggtgcagc 1140
 cacgtctgct ccgcacgcc ggtggagacg gtgaacatca gcaatgagct catctcctac 1200
 aaccaagaca cggctttcca caaggaaac gcagctgcct acatccacat gatgcccaac 1260
 gccattcccc ccggggaccg ggaggtggac aacgatgagg aggaggagag tcctttcgat 1320
 cgcattgccc agatctatca gacatcccca gatggtgacc atgttgaga gtctgtctgg 1380
 gagctggact gcgaggggga gatttcttta gacaaaataa cacctttcac cccaaatgga 1440
 ttccattaaa ctgcattaag aaacccctc atggatctgc ataaccacac agacattgac 1500
 aagcatgcac acacaagcaa atacatggct ttcca 1535

<210> 15
 <211> 480
 <212> PRT
 <213> Homo sapiens

<400> 15
 Met Leu Pro Pro Arg Ser Asn Asp Thr Ala Tyr Pro Gly Gln Leu Ala
 1 5 10 15
 Leu Tyr Gln Gln Leu Ala Gln Gly Asn Ala Val Gly Gly Ser Ala Gly
 20 25 30
 Ala Pro Pro Leu Gly Pro Val Gln Val Val Thr Ala Cys Leu Leu Thr
 35 40 45

Leu Leu Val Ile Trp Thr Leu Leu Gly Asn Val Leu Val Ser Ala Ala
 50 55 60
 Ile Val Trp Ser Arg His Leu Arg Ala Lys Met Thr Asn Val Phe Ile
 65 70 75 80
 Val Ser Leu Pro Val Ser Asp Leu Phe Val Ala Leu Leu Val Met Ser
 85 90 95
 Trp Lys Ala Val Ala Glu Val Ala Gly Tyr Trp Pro Phe Glu Ala Phe
 100 105 110
 Cys Asp Val Trp Val Ala Phe Asp Ile Met Cys Ser Thr Ala Ser Ile
 115 120 125
 Leu Asn Leu Cys Val Ile Ser Val Ala Arg Tyr Trp Ala Ile Ser Arg
 130 135 140
 Pro Phe Arg Tyr Glu Arg Lys Met Thr Gln Arg Met Ala Leu Val Met
 145 150 155 160
 Val Arg Pro Ala Trp Thr Leu Ser Ser Leu Ile Ser Phe Ile Pro Val
 165 170 175
 Gln Leu Asn Trp His Arg Asp Gln Ala Val Ser Trp Gly Gly Leu Asp
 180 185 190
 Leu Pro Asn Asn Leu Ala Asn Trp Thr Pro Trp Glu Glu Ala Val Trp
 195 200 205
 Glu Pro Asp Val Arg Ala Glu Asn Cys Asp Ser Ser Leu Asn Arg Thr
 210 215 220
 Tyr Ala Ile Pro Ser Ser Leu Ile Ser Phe Tyr Ile Pro Met Ala Ile
 225 230 235 240
 Met Ile Val Thr Tyr Thr Arg Ile Tyr Arg Ile Ala Gln Val Gln Ile
 245 250 255
 Arg Arg Ile Ser Ser Leu Glu Arg Ala Ala Glu His Val Gln Ser Cys
 260 265 270
 Arg Ser Ser Ala Gly Cys Thr Pro Asp Thr Ser Leu Arg Phe Ser Ile
 275 280 285
 Lys Lys Glu Thr Glu Val Leu Lys Thr Leu Ser Val Ile Met Gly Val
 290 295 300
 Phe Val Cys Cys Trp Leu Pro Phe Phe Ile Leu Asn Cys Met Val Pro
 305 310 315 320
 Phe Cys Ser Gly His Pro Lys Gly Pro Pro Ala Gly Phe Pro Cys Val
 325 330 335
 Ser Glu Thr Thr Phe Asp Val Phe Ile Trp Phe Cys Trp Ala Asn Ser
 340 345 350

Ser Leu Asn Pro Val Pro Ser His Tyr Ala Phe Asn Ala Asp Phe Arg
 355 360 365
 Lys Val Phe Ala Gln Leu Leu Gly Cys Ser His Val Cys Ser Arg Thr
 370 375 380
 Pro Val Glu Thr Val Asn Ile Ser Asn Glu Leu Ile Ser Tyr Asn Gln
 385 390 395 400
 Asp Thr Val Phe His Lys Glu Ile Ala Ala Ala Tyr Ile His Met Met
 405 410 415
 Pro Asn Ala Ile Pro Pro Gly Asp Arg Glu Val Asp Asn Asp Glu Glu
 420 425 430
 Glu Glu Ser Pro Phe Asp Arg Met Ser Gln Ile Tyr Gln Thr Ser Pro
 435 440 445
 Asp Gly Asp His Val Ala Glu Ser Val Trp Glu Leu Asp Cys Glu Gly
 450 455 460
 Glu Ile Ser Leu Asp Lys Ile Thr Pro Phe Thr Pro Asn Gly Phe His
 465 470 475 480

<210> 16
 <211> 2657
 <212> DNA
 <213> Homo sapiens

<400> 16
 gattcatgaa gatgttgaca agactacaag ttcttatgtt agctttgttt tcaaagggat 60
 ttttagtctc tttaggagat cacaacttta tgaggagaga aattaaaata gaaggagacc 120
 ttgttttagg gggcttattt cctattaatg aaaaaggcac tgggaactgaa gagtgtggac 180
 gaatcaatga agacagaggt atccaacgcc tggaggccat gttgtttgcc attgatgaaa 240
 tcaacaaaga caattacttg cttccaggag tgaagctggg gggtcacatt ttggatacat 300
 gttcaagaga cacttatgca ttagagcagt cactggagtt tgtcagagca tcgttgacta 360
 aagtggatga agctgaatat attggcgggt cgtacagcag tgctattcaa gaaaacatcc 420
 cactactcat tgcaggagtc attggcgggt cgtacagcag tgctattcaa gaaaacatcc 480
 acctgctgag gctcttccag atccctcaga taagctacgc ctccaccagt gccaaactca 540
 gcgacaaatc gcgctatgat tattttgcca ggaccgtgcc ccctgacttc taccaggcca 600
 aagccatggc cgagatcttg cgctacttta actggacctt tgtgtccact gttgcctctg 660
 aaggtgacta tggggagaca gggattgagg ccttcgagca ggaagcaagg ctacgcaaca 720
 tctgcatcgc cactgctgaa aaggtggggc gctccaacat ccgcaagtcc tacgacagcg 780
 tgatccgtga gctcctgcag aaacctaacg cgcgagttgt ggctcctgtt atgcgcagtg 840
 atgactcacg agagttgatc gctgcagcca gccgcgtgaa tgcttccttc acctgggtgg 900
 ccagcgatgg ctgggggtgca caggagagca ttgtcaaggg cagtgcagcac gtcgcctatg 960
 gagccatcac cctggagctg gcgtcccacc ctgttcgtca gtttgatcgc tacttccaga 1020
 gcctcaaccc ctacaacaat catcgtaacc cctgggtccg agacttcttg gagcagaagt 1080
 tccagtgcag cctccagaac aagagaaacc acagacagat ttgtgacaag cacctggcca 1140
 ttgacagcag caactatgaa caagaatcca agatcatgtt tgtggtgaat gcagtgtatg 1200
 ccatggcgca tgcgctgcac aaaatgcaac gcaccctctg tcccaacacc accaagctct 1260
 gtgatgcaat gaagatcctg gatggaaaga agttgtacaa agattatttg ctgaaaatca 1320
 acttccttgc tccattcaac ccaaataaag gagcagacag cattgtgaag tttgacactt 1380


```

acggagacgg gatgggaaga tacaacgtgt tcaacttcca gcatataggt ggaaagtatt 1440
cctacttaaa agttggccac tgggcagaaa ctttatatct agatgtggac tctattcatt 1500
gggtcccggaa ctccagtcctc acttcccagt gcagtgatcc ctgtgcccc aatgaaatga 1560
aaaacatgca gccaggagat gtttgctgct ggatctgcat cccatgtgag ccctatgaat 1620
acctggttga tgagttcacc tgcattggatt gtggccctgg ccagtggccc actgcagacc 1680
tatctggatg ctacaacctt ccagaggatt acatcagggtg ggaagatgcc tgggcaatag 1740
gcccagtcac tattgcctgc ctgggtttta tgtgtacatg catagtcata actgttttta 1800
tcaagcacia caacacaccc ttgggtcaaag catcaggccg agaactctgc tacatcttgt 1860
tatttgaggt tagcctgtcc tattgcatga cattcttctt cattgctaag ccacgcctg 1920
tcactctgtc attgcccga cttgggcttg ggacctcctt tgccatctgt tattcagctc 1980
tcctgaccaa gacaaactgc atcgctcgca tctttgatgg ggtcaagaat ggcgctcaga 2040
ggccaaaatt catcagcccc agttctcagg tttttatctg cctgggtttg atactgggtg 2100
aaattgtgat ggtgtctgtg tggcttatct tggagactcc aggtactaga agatacacc 2160
tgccagagaa gcgggaaaca gtcaccta aatgcaatgt caaagattcc agcatgttga 2220
tctctctgac ctatgacgtg gttctggtga ttctatgcac tgtgtatgcc ttcaaaacaa 2280
ggaagtgtcc tgaaaacttc aatgaagcca agttcatagg cttcaccatg tacaccacct 2340
gcatcatctg gttggcattc ctccctatat tttatgtgac atcaagtgcac tacagagtac 2400
agacgacaac aatgtgcac tccgttagct tgagtgggtt cgtgggtctg ggctgtttgt 2460
ttgcccccaa ggtgcacatt gtccgtgtcc aaccacagaa gaatgtggtc acacacagac 2520
ttcacctcaa cagggttcagt gtcagtggaa ctgcgaccac atattctcag gcctctgcaa 2580
gcacgtatgt gccaacggtg tgcaatgggc ggggaagtcct cgactccacc acctcatctc 2640
tgtgattgtg aattgca 2657

```

<210> 17

<211> 879

<212> PRT

<213> Homo sapiens

<400> 17

```

Met Lys Met Leu Thr Arg Leu Gln Val Leu Met Leu Ala Leu Phe Ser
 1              5              10              15

Lys Gly Phe Leu Val Ser Leu Gly Asp His Asn Phe Met Arg Arg Glu
              20              25              30

Ile Lys Ile Glu Gly Asp Leu Val Leu Gly Gly Leu Phe Pro Ile Asn
 35              40              45

Glu Lys Gly Thr Gly Thr Glu Glu Cys Gly Arg Ile Asn Glu Asp Arg
 50              55              60

Gly Ile Gln Arg Leu Glu Ala Met Leu Phe Ala Ile Asp Glu Ile Asn
 65              70              75              80

Lys Asp Asn Tyr Leu Leu Pro Gly Val Lys Leu Gly Val His Ile Leu
      85              90              95

Asp Thr Cys Ser Arg Asp Thr Tyr Ala Leu Glu Gln Ser Leu Glu Phe
 100              105              110

Val Arg Ala Ser Leu Thr Lys Val Asp Glu Ala Glu Tyr Met Cys Pro
 115              120              125

Asp Gly Ser Tyr Ala Ile Gln Glu Asn Ile Pro Leu Leu Ile Ala Gly
 130              135              140

```

Val	Ile	Gly	Gly	Ser	Tyr	Ser	Ser	Val	Ser	Ile	Gln	Val	Ala	Asn	Leu	145	150	155	160
Leu	Arg	Leu	Phe	Gln	Ile	Pro	Gln	Ile	Ser	Tyr	Ala	Ser	Thr	Ser	Ala	165	170	175	
Lys	Leu	Ser	Asp	Lys	Ser	Arg	Tyr	Asp	Tyr	Phe	Ala	Arg	Thr	Val	Pro	180	185	190	
Pro	Asp	Phe	Tyr	Gln	Ala	Lys	Ala	Met	Ala	Glu	Ile	Leu	Arg	Tyr	Phe	195	200	205	
Asn	Trp	Thr	Tyr	Val	Ser	Thr	Val	Ala	Ser	Glu	Gly	Asp	Tyr	Gly	Glu	210	215	220	
Thr	Gly	Ile	Glu	Ala	Phe	Glu	Gln	Glu	Ala	Arg	Leu	Arg	Asn	Ile	Cys	225	230	235	240
Ile	Ala	Thr	Ala	Glu	Lys	Val	Gly	Arg	Ser	Asn	Ile	Arg	Lys	Ser	Tyr	245	250	255	
Asp	Ser	Val	Ile	Arg	Glu	Leu	Leu	Gln	Lys	Pro	Asn	Ala	Arg	Val	Val	260	265	270	
Val	Leu	Phe	Met	Arg	Ser	Asp	Asp	Ser	Arg	Glu	Leu	Ile	Ala	Ala	Ala	275	280	285	
Ser	Arg	Val	Asn	Ala	Ser	Phe	Thr	Trp	Val	Ala	Ser	Asp	Gly	Trp	Gly	290	295	300	
Ala	Gln	Glu	Ser	Ile	Val	Lys	Gly	Ser	Glu	His	Val	Ala	Tyr	Gly	Ala	305	310	315	320
Ile	Thr	Leu	Glu	Leu	Ala	Ser	His	Pro	Val	Arg	Gln	Phe	Asp	Arg	Tyr	325	330	335	
Phe	Gln	Ser	Leu	Asn	Pro	Tyr	Asn	Asn	His	Arg	Asn	Pro	Trp	Phe	Arg	340	345	350	
Asp	Phe	Trp	Glu	Gln	Lys	Phe	Gln	Cys	Ser	Leu	Gln	Asn	Lys	Arg	Asn	355	360	365	
His	Arg	Gln	Ile	Cys	Asp	Lys	His	Leu	Ala	Ile	Asp	Ser	Ser	Asn	Tyr	370	375	380	
Glu	Gln	Glu	Ser	Lys	Ile	Met	Phe	Val	Val	Asn	Ala	Val	Tyr	Ala	Met	385	390	395	400
Ala	His	Ala	Leu	His	Lys	Met	Gln	Arg	Thr	Leu	Cys	Pro	Asn	Thr	Thr	405	410	415	
Lys	Leu	Cys	Asp	Ala	Met	Lys	Ile	Leu	Asp	Gly	Lys	Lys	Leu	Tyr	Lys	420	425	430	
Asp	Tyr	Leu	Leu	Lys	Ile	Asn	Phe	Leu	Ala	Pro	Phe	Asn	Pro	Asn	Lys	435	440	445	

Gly Ala Asp Ser Ile Val Lys Phe Asp Thr Tyr Gly Asp Gly Met Gly
 450 455 460
 Arg Tyr Asn Val Phe Asn Phe Gln His Ile Gly Gly Lys Tyr Ser Tyr
 465 470 475 480
 Leu Lys Val Gly His Trp Ala Glu Thr Leu Tyr Leu Asp Val Asp Ser
 485 490 495
 Ile His Trp Ser Arg Asn Ser Val Pro Thr Ser Gln Cys Ser Asp Pro
 500 505 510
 Cys Ala Pro Asn Glu Met Lys Asn Met Gln Pro Gly Asp Val Cys Cys
 515 520 525
 Trp Ile Cys Ile Pro Cys Glu Pro Tyr Glu Tyr Leu Val Asp Glu Phe
 530 535 540
 Thr Cys Met Asp Cys Gly Pro Gly Gln Trp Pro Thr Ala Asp Leu Ser
 545 550 555 560
 Gly Cys Tyr Asn Leu Pro Glu Asp Tyr Ile Arg Trp Glu Asp Ala Trp
 565 570 575
 Ala Ile Gly Pro Val Thr Ile Ala Cys Leu Gly Phe Met Cys Thr Cys
 580 585 590
 Ile Val Ile Thr Val Phe Ile Lys His Asn Asn Thr Pro Leu Val Lys
 595 600 605
 Ala Ser Gly Arg Glu Leu Cys Tyr Ile Leu Leu Phe Gly Val Ser Leu
 610 615 620
 Ser Tyr Cys Met Thr Phe Phe Phe Ile Ala Lys Pro Ser Pro Val Ile
 625 630 635 640
 Cys Ala Leu Arg Arg Leu Gly Leu Gly Thr Ser Phe Ala Ile Cys Tyr
 645 650 655
 Ser Ala Leu Leu Thr Lys Thr Asn Cys Ile Ala Arg Ile Phe Asp Gly
 660 665 670
 Val Lys Asn Gly Ala Gln Arg Pro Lys Phe Ile Ser Pro Ser Ser Gln
 675 680 685
 Val Phe Ile Cys Leu Gly Leu Ile Leu Val Gln Ile Val Met Val Ser
 690 695 700
 Val Trp Leu Ile Leu Glu Thr Pro Gly Thr Arg Arg Tyr Thr Leu Pro
 705 710 715 720
 Glu Lys Arg Glu Thr Val Ile Leu Lys Cys Asn Val Lys Asp Ser Ser
 725 730 735
 Met Leu Ile Ser Leu Thr Tyr Asp Val Val Leu Val Ile Leu Cys Thr
 740 745 750

Val Tyr Ala Phe Lys Thr Arg Lys Cys Pro Glu Asn Phe Asn Glu Ala
 755 760 765
 Lys Phe Ile Gly Phe Thr Met Tyr Thr Thr Cys Ile Ile Trp Leu Ala
 770 775 780
 Phe Leu Pro Ile Phe Tyr Val Thr Ser Ser Asp Tyr Arg Val Gln Thr
 785 790 795 800
 Thr Thr Met Cys Ile Ser Val Ser Leu Ser Gly Phe Val Val Leu Gly
 805 810 815
 Cys Leu Phe Ala Pro Lys Val His Ile Val Leu Phe Gln Pro Gln Lys
 820 825 830
 Asn Val Val Thr His Arg Leu His Leu Asn Arg Phe Ser Val Ser Gly
 835 840 845
 Thr Ala Thr Thr Tyr Ser Gln Ala Ser Ala Ser Thr Tyr Val Pro Thr
 850 855 860
 Val Cys Asn Gly Arg Glu Val Leu Asp Ser Thr Thr Ser Ser Leu
 865 870 875

<210> 18
 <211> 1366
 <212> DNA
 <213> Homo sapiens

<400> 18
 atgggtctgg ccatggagca cggaggggtcc tacgctcggg cgggggggcag ctctcggggc 60
 tgctgggtatt acctgcgcta cttcttcctc ttcgtctccc tcatccaatt cctcatcatc 120
 ctggggctcg tgctcttcat ggtctatggc aacgtgcacg tgagcacaga gtccaacctg 180
 caggccaccg agcgccgagc cgagggccta tacagtcagc tcctagggct cacggcctcc 240
 cagtccaact tgaccaagga gctcaacttc accaccgcg ccaaggatgc catcatgcag 300
 atgtggctga atgctcgccg cgacctggac cgcataaatg ccagcttcg ccagtgccag 360
 ggtgaccggg taatctacac gaacaatcag aggtacatgg ctgccatcat cttgagtgag 420
 aagcaatgca gagatcaatt caaggacatg aacaagagct gcgatgcctt gctcttcatg 480
 ctgaatcaga aggtgaagac gctggagggtg gagatagcca aggagaagac catttgact 540
 aaggataagg aaagcgtgct gctgaacaaa cgcgtggcgg aggaacagct ggttgaatgc 600
 gtgaaaaccc gggagctgca gcaccaagag cgccagctgg ccaaggagca actgcaaaag 660
 gtgcaagccc tctgcctgcc cctggacaag gacaagtttg agatggacct tcgtaacctg 720
 tggagggact ccattatccc acgcagcctg gacaacctgg gttacaacct ctaccatccc 780
 ctgggctcgg aattggcctc catccgcaga gcctgcgacc acatgcccag cctcatgagc 840
 tccaaggtgg aaggctcagt cgggagcctc cgggcggata tcgaacgcgt ggcccgcgag 900
 aactcagacc tccaacgcca gaagctggaa gcccagcagg gcctgcgggc cagtcaggag 960
 gcgaaacaga aggtggagaa ggaggctcag gcccgggagg ccaagctcca agctgaatgc 1020
 tcccggcaga cccagctagc gctggaggag aaggcggtgc tgcggaagga acgagacaac 1080
 ctggccaagg agctggaaga gaagaagagg gaggcggagc agctcaggat ggagctggcc 1140
 atcagaaact cagccctgga cacctgcata aagaccaagt cgcagccgat gatgccagt 1200
 tcaaggccca tgggccctgt ccccaacccc cagcccatcg accagctag cctggaggag 1260
 ttcaagagga agatcctgga gtcccagagg cccctgcag gcatacctgt agccccatcc 1320
 agtggctgag gaggtccgg cactgaccta agggcgaatc ccagca 1366

<210> 19

<400> 19

Asp His Met Pro Ser Leu Met Ser Ser Lys Val Glu Gly Gln Cys Arg

275	280	285
Ser Leu Arg Ala Asp Ile Glu Arg Val Ala Arg Glu Asn Ser Asp Leu		
290	295	300
Gln Arg Gln Lys Leu Glu Ala Gln Gln Gly Leu Arg Ala Ser Gln Glu		
305	310	315
Ala Lys Gln Lys Val Glu Lys Glu Ala Gln Ala Arg Glu Ala Lys Leu		
	325	330
Gln Ala Glu Cys Ser Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala		
	340	350
Val Leu Arg Lys Glu Arg Asp Asn Leu Ala Lys Glu Leu Glu Glu Lys		
	355	365
Lys Arg Glu Ala Glu Gln Leu Arg Met Glu Leu Ala Ile Arg Asn Ser		
	370	380
Ala Leu Asp Thr Cys Ile Lys Thr Lys Ser Gln Pro Met Met Pro Val		
385	390	395
Ser Arg Pro Met Gly Pro Val Pro Asn Pro Gln Pro Ile Asp Pro Ala		
	405	410
Ser Leu Glu Glu Phe Lys Arg Lys Ile Leu Glu Ser Gln Arg Pro Pro		
	420	430
Ala Gly Ile Pro Val Ala Pro Ser Ser Gly		
	435	440

<210> 20
 <211> 1421
 <212> DNA
 <213> Homo sapiens

<400> 20
 gaattctagg tgggtggtgag cagggacggt gcaccggacg gcgggatcga gcaaattgggt 60
 ctggccatgg agtacggagg gtcctacgct cgggcggggg gcagctctcg gggctgctgg 120
 tattacctgc gctacttctt cctcttcgtc tccctcatcc aattcctcat catcctgggg 180
 ctctgtctct tcatggtcta tggcgacgtg cacgtgagca cagagtccaa cctgcaggcc 240
 accgagcgcc gagccgaggg cctatacagt cagctcctag ggctcacggc ctcccagtc 300
 aacttgacca aggagctcaa cttcaccacc cgcgccaagg atgccatcat gcagatgtgg 360
 ctgaatgctc gccgcgacct ggaccgcatc aatgccagct tccgccagtg ccagggtgac 420
 cgggtcatct acacgaacaa tcagagggtac atggctgcca tcatcttgag tgagaagcaa 480
 tgcagagatc aattcaagga catgaacaag agctgcatg ccttgctctt catgctgaat 540
 cagaagggtga agacgctgga ggtggagata gccaaaggaga agaccatttg cactaaggat 600
 aaggaaagcg tgctgctgaa caaacgcgtg gcggagggaac agctggttga atgcgtgaaa 660
 acccgggagc tgcagcacca agagcgccag ctggccaagg agcaactgca aaagggtgcaa 720
 gccctctgcc tgcccctgga caaggacaag tttgagatgg accttcgtaa cctgtggagg 780
 gactccatta tcccacgcag cctggacaac ctgggttaca acctctacca tcccctgggc 840
 tcggaattgg cctccatccg cagagcctgc gaccacatgc ccagcctcat gagctccaag 900
 gtggaggagc tggcccgag cctccgggag gatatcgaac gcgtggcccg cgagaactca 960
 gacctccaac gccagaagct ggaagcccag cagggcctgc gggccagtca ggaggcgaaa 1020
 cagaagggtg agaaggaggc tcaggcccgg gaggccaagc tccaagctga atgctcccgg 1080

cagacccagc tagcgctgga ggagaaggcg gtgctgcgga aggaacgaga caacctggcc 1140
aaggagctgg aagagaagaa gagggaggcg gagcagctca ggatggagct ggccatcaga 1200
aactcagccc tggacacctg catcaagacc aagtcgcagc cgatgatgcc agtgtcaagg 1260
cccatggggc ctgtcccca cccccagccc atcgacccag ctagcctgga ggagttcaag 1320
aggaagatcc tggagtccca gagggccctt gcaggcatcc ctgtagcccc atccagtggc 1380
tgaggaggct ccaggcctga ggaccaaggg atggcccgac t 1421

<210> 21
<211> 442
<212> PRT
<213> Homo sapiens

<400> 21

Met	Gly	Leu	Ala	Met	Glu	Tyr	Gly	Gly	Ser	Tyr	Ala	Arg	Ala	Gly	Gly	1	5	10	15
Ser	Ser	Arg	Gly	Cys	Trp	Tyr	Tyr	Leu	Arg	Tyr	Phe	Phe	Leu	Phe	Val	20	25	30	
Ser	Leu	Ile	Gln	Phe	Leu	Ile	Ile	Leu	Gly	Leu	Val	Leu	Phe	Met	Val	35	40	45	
Tyr	Gly	Asp	Val	His	Val	Ser	Thr	Glu	Ser	Asn	Leu	Gln	Ala	Thr	Glu	50	55	60	
Arg	Arg	Ala	Glu	Gly	Leu	Tyr	Ser	Gln	Leu	Leu	Gly	Leu	Thr	Ala	Ser	65	70	75	80
Gln	Ser	Asn	Leu	Thr	Lys	Glu	Leu	Asn	Phe	Thr	Thr	Arg	Ala	Lys	Asp	85	90	95	
Ala	Ile	Met	Gln	Met	Trp	Leu	Asn	Ala	Arg	Arg	Asp	Leu	Asp	Arg	Ile	100	105	110	
Asn	Ala	Ser	Phe	Arg	Gln	Cys	Gln	Gly	Asp	Arg	Val	Ile	Tyr	Thr	Asn	115	120	125	
Asn	Gln	Arg	Tyr	Met	Ala	Ala	Ile	Ile	Leu	Ser	Glu	Lys	Gln	Cys	Arg	130	135	140	
Asp	Gln	Phe	Lys	Asp	Met	Asn	Lys	Ser	Cys	Asp	Ala	Leu	Leu	Phe	Met	145	150	155	160
Leu	Asn	Gln	Lys	Val	Lys	Thr	Leu	Glu	Val	Glu	Ile	Ala	Lys	Glu	Lys	165	170	175	
Thr	Ile	Cys	Thr	Lys	Asp	Lys	Glu	Ser	Val	Leu	Leu	Asn	Lys	Arg	Val	180	185	190	
Ala	Glu	Glu	Gln	Leu	Val	Glu	Cys	Val	Lys	Thr	Arg	Glu	Leu	Gln	His	195	200	205	
Gln	Glu	Arg	Gln	Leu	Ala	Lys	Glu	Gln	Leu	Gln	Lys	Val	Gln	Ala	Leu	210	215	220	
Cys	Leu	Pro	Leu	Asp	Lys	Asp	Lys	Phe	Glu	Met	Asp	Leu	Arg	Asn	Leu				

225		230		235		240
Trp Arg Asp Ser Ile Ile Pro Arg Ser Leu Asp Asn Leu Gly Tyr Asn						
	245		250		255	
Leu Tyr His Pro Leu Gly Ser Glu Leu Ala Ser Ile Arg Arg Ala Cys						
	260		265		270	
Asp His Met Pro Ser Leu Met Ser Ser Lys Val Glu Glu Leu Ala Arg						
	275		280		285	
Ser Leu Arg Ala Asp Ile Glu Arg Val Ala Arg Glu Asn Ser Asp Leu						
	290		295		300	
Gln Arg Gln Lys Leu Glu Ala Gln Gln Gly Leu Arg Ala Ser Gln Glu						
305		310		315		320
Ala Lys Gln Lys Val Glu Lys Glu Ala Gln Ala Arg Glu Ala Lys Leu						
	325		330		335	
Gln Ala Glu Cys Ser Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala						
	340		345		350	
Val Leu Arg Lys Glu Arg Asp Asn Leu Ala Lys Glu Leu Glu Glu Lys						
	355		360		365	
Lys Arg Glu Ala Glu Gln Leu Arg Met Glu Leu Ala Ile Arg Asn Ser						
	370		375		380	
Ala Leu Asp Thr Cys Ile Lys Thr Lys Ser Gln Pro Met Met Pro Val						
385		390		395		400
Ser Arg Pro Met Gly Pro Val Pro Asn Pro Gln Pro Ile Asp Pro Ala						
	405		410		415	
Ser Leu Glu Glu Phe Lys Arg Lys Ile Leu Glu Ser Gln Arg Pro Pro						
	420		425		430	
Ala Gly Ile Pro Val Ala Pro Ser Ser Gly						
	435		440			

<210> 22
 <211> 2024
 <212> DNA
 <213> Homo sapiens

<400> 22
 atgggtctgg ccatggagca cggagggtcc tacgctcggg cggggggcag ctctcggggc 60
 tgctggtatt acctgcgcta cttcttcttc ttctgtctccc tcatccaatt cctcatcatc 120
 ctggggctcg tgctcttcat ggtctatggc aacgtgcacg tgagcacaga gtccaacctg 180
 caggccaccg agcgccgagc cgagggccta tacagtcagc tcctagggct cacggcctcc 240
 cagtccaact tgaccaagga gctcaacttc accaccgcg ccaaggatgc catcatgcag 300
 atgtggctga atgctcgtcg cgacctggac cgcataatg ccagcttccg ccagtgccag 360
 ggtgaccggg tcatctacac gaacaatcag aggtacatgg ctgccatcat cttgagttag 420
 aagcaatgca gagatcaatt caaggacatg aacaagagct gcgatgcctt gctcttcatg 480
 ctgaatcaga aggtgaagac gctggagggtg gagatagcca aggagaagac catttgact 540


```

aaggataagg aaagcgtgct gctgaacaaa cgcgtggcgg aggaacagct ggttgaatgc 600
gtgaaaaccc gggagctgca gcaccaagag cgccagctgg ccaaggagca actgcaaagg 660
gtgcaagccc tctgcttccc cctggacaag gacaagtttg agatggacct tcgtaacctg 720
tggagggact ccattatccc acgcagcctg gacaacctgg gttacaacct ctaccatccc 780
ctgggctcgg aattggcctc catccgcaga gcctgcgacc acatgcccag cctcgtgagc 840
tccaaggtgg aggagctggc ccggagcctc cgggcggata tcgaacgcgt ggcccgcgag 900
aactcagacc tccaacgcca gaagctggaa gcccagcagg gcctgcgggc cagtcaggag 960
cgaacacaga aggtggagaa ggaggctcag gcccgggagg ccaagctcca agctgaatgc 1020
tcccggcaga ccagctagc gctggaggag aaggcgggtg tcggaagga acgagacaac 1080
ctggccaagg agctggaaga gaagaagagg gaggcggagc agctcaggat ggagctggcc 1140
atcagaaact cagccctgga cacctgcatc aagaccaagt cgcagccgat gatgccagt 1200
tcaaggccca tggggccctgt cccaacccc cagcccatcg accagctag cctggaggag 1260
ttcaagagga agatccctgga gtcccagagg cccctgcag gcatccctgt agcccatcc 1320
agtggctgag gaggtctccag gcctgaggag caagggatgg cccgactcgg cggtttgagg 1380
aggatgcagg gatatgctca cagcgcccga cacaacccc tccgcccggc cccaaccacc 1440
cagggccacc atcagacaac tccctgcatg caaaccccta gtaccctctc acaccgcac 1500
ccgcgctcga tgatccctca ccagagcac acggccgcgg agatgacgtc acgcaagcaa 1560
cggcgctgac gtcacatata accgtggtga tggcgctcag tggccatgta gacgtcacga 1620
agagatatag cgatggcgtc gtgcagatgc agcacgtcgc acacagacat ggggaacttg 1680
gcatgacgtc acaccgagat gcagcaacga cgtcacgggc catgtcgacg tcacacatat 1740
taatgtcaca cagacgcggc gatggcatca cacagacggg gatgatgtca cacacagaca 1800
cagtgacaac acacaccatg acaacgacac ctatagatat ggcaccaaca tcacatgcac 1860
gcatgccctt tcacacacac tttctaccca attctcacct agtgtcacgt tcccccgacc 1920
ctggcacacg ggccaaggta cccacaggat cccatcccct cccgcacagc cctgggcccc 1980
agcacctccc ctctccagc ctctggcct cccggtagta cacg 2024

```

<210> 23
 <211> 442
 <212> PRT
 <213> Homo sapiens

<400> 23

```

Met Gly Leu Ala Met Glu His Gly Gly Ser Tyr Ala Arg Ala Gly Gly
  1              5              10              15

Ser Ser Arg Gly Cys Trp Tyr Tyr Leu Arg Tyr Phe Phe Leu Phe Val
  20              25              30

Ser Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Val
  35              40              45

Tyr Gly Asn Val His Val Ser Thr Glu Ser Asn Leu Gln Ala Thr Glu
  50              55              60

Arg Arg Ala Glu Gly Leu Tyr Ser Gln Leu Leu Gly Leu Thr Ala Ser
  65              70              75              80

Gln Ser Asn Leu Thr Lys Glu Leu Asn Phe Thr Thr Arg Ala Lys Asp
  85              90              95

Ala Ile Met Gln Met Trp Leu Asn Ala Arg Arg Asp Leu Asp Arg Ile
 100              105              110

Asn Ala Ser Phe Arg Gln Cys Gln Gly Asp Arg Val Ile Tyr Thr Asn
 115              120              125

```

Asn Gln Arg Tyr Met Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Arg
 130 135 140

Asp Gln Phe Lys Asp Met Asn Lys Ser Cys Asp Ala Leu Leu Phe Met
 145 150 155 160

Leu Asn Gln Lys Val Lys Thr Leu Glu Val Glu Ile Ala Lys Glu Lys
 165 170 175

Thr Ile Cys Thr Lys Asp Lys Glu Ser Val Leu Leu Asn Lys Arg Val
 180 185 190

Ala Glu Glu Gln Leu Val Glu Cys Val Lys Thr Arg Glu Leu Gln His
 195 200 205

Gln Glu Arg Gln Leu Ala Lys Glu Gln Leu Gln Arg Val Gln Ala Leu
 210 215 220

Cys Leu Pro Leu Asp Lys Asp Lys Phe Glu Met Asp Leu Arg Asn Leu
 225 230 235 240

Trp Arg Asp Ser Ile Ile Pro Arg Ser Leu Asp Asn Leu Gly Tyr Asn
 245 250 255

Leu Tyr His Pro Leu Gly Ser Glu Leu Ala Ser Ile Arg Arg Ala Cys
 260 265 270

Asp His Met Pro Ser Leu Val Ser Ser Lys Val Glu Glu Leu Ala Arg
 275 280 285

Ser Leu Arg Ala Asp Ile Glu Arg Val Ala Arg Glu Asn Ser Asp Leu
 290 295 300

Gln Arg Gln Lys Leu Glu Ala Gln Gln Gly Leu Arg Ala Ser Gln Glu
 305 310 315 320

Ala Lys Gln Lys Val Glu Lys Glu Ala Gln Ala Arg Glu Ala Lys Leu
 325 330 335

Gln Ala Glu Cys Ser Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala
 340 345 350

Val Leu Arg Lys Glu Arg Asp Asn Leu Ala Lys Glu Leu Glu Glu Lys
 355 360 365

Lys Arg Glu Ala Glu Gln Leu Arg Met Glu Leu Ala Ile Arg Asn Ser
 370 375 380

Ala Leu Asp Thr Cys Ile Lys Thr Lys Ser Gln Pro Met Met Pro Val
 385 390 395 400

Ser Arg Pro Met Gly Pro Val Pro Asn Pro Gln Pro Ile Asp Pro Ala
 405 410 415

Ser Leu Glu Glu Phe Lys Arg Lys Ile Leu Glu Ser Gln Arg Pro Pro
 420 425 430

Ala Gly Ile Pro Val Ala Pro Ser Ser Gly
 435 440

<210> 24
 <211> 8640
 <212> DNA
 <213> Homo sapiens

<400> 24

```

agctgatgat ggccagggac cccaggggac gtggggccct gtgggggtctg gccccagga 60
gcaagacctc tgatgatgct ggtgtctggg agtgagcacc atgcccacatca cccaggacaa 120
tgccgtgctg cacctgcccc tcctctacca gtggctgcag aacagcctgc aggaaggtgg 180
ggatgggccc gagcagcggc tctgccaggc ggccatccag aagctgcagg agtacatcca 240
gctgaacttt gctgtggatg agagtacggt cccacctgat cacagcccc ccgaaatgga 300
gatctgtact gtgtacctca ccaaggagct gggggacaca gagactgtgg gcctgagttt 360
tggaacatc cctgttttcg gggactatgg tgaaaagcgc agggggggca agaagaggaa 420
aaccaccag ggtcctgtgc tggatgtggg ctgcatctgg gtgacagagc tgaggaagaa 480
cagcccagca ggaagagtg ggaaggtccg actgcgggat gagatcctct cactgaatgg 540
gcagctgatg gttggagttg atgtcagtgg ggccagttac ctggctgagc agtgctggaa 600
tgccggtctt atctacctga tcatgtctgc tcgctttaag cacaagccc actccactta 660
taatggcaac agtagcaaca gctctgaacc aggagaaaca cctaccttgg agctgggtga 720
ccgaactgcg aaaaagggga aacgaaccag aaagtttggg gtcacttcca ggctcctgc 780
caacaaggcc cctgaagaat ccaagggcag cgctggctgt gaggtgtcca gtgaccccag 840
cactgagctg gagaacggcc tggaccctga acttggaaac ggccatgtct ttcagctaga 900
aaatggccca gattctctca aggaggtggc tggaccccat ctagagaggt cagaagtgga 960
cagagggaca gagcatagaa ttccaaagac agatgtcct ctgaccacaa gcaatgacaa 1020
acgccgttc tcaaaagggtg ggaagacgga cttccaatcg agtgactgcc tggcacggtc 1080
caaggaggaa gttggccgaa tatggaagat ggagctgctc aaagaatcgg atgggctggg 1140
aattcaggtt agtggaggcc gaggatcaaa gcgtcacct cacgctatcg ttgtcactca 1200
agtgaaggaa ggaggtgccg ctcacaggct cagggatggc aggtgtcct taggagatga 1260
gctgctggta atcaatggtc atttactggt cgggctctcc cacgaggaag cagtggccat 1320
tcttcgctcc gccacgggaa tgggtgcagct tgtgggtggc agcaaggtag gtgtgctttc 1380
tgcatctcag atgcttggga cagatgaacc ccaagatgtg tgcggtgctg aggaatccaa 1440
ggggaacttg gaaagtccea aacagggcag caataaaatc aagctcaaga gtcgcctttc 1500
aggtaggtgg gggctctacc tgatgcagcc tgtcgggggt gtacaccgcc ttgagtcagt 1560
tgaagaatat aacgagctga tgggtgcgaa tggggacccc cgcatccgga tgttgagggt 1620
ctcccagat ggccggaaac actccctccc gcagctgctg gactcttcca gtgcctcaca 1680
ggaataccac attgtgaaga agtctacccg ctcttaagc acgactcagg tggaatctcc 1740
ttggaggctc attcggccat ccgtcatctc gatcttggg ttgtacaaag aaaaaggcaa 1800
gggccttggc tttagtattg ctggaggtcg agactgcatt cgtggacaga tggggatttt 1860
tgtcaagacc atcttcccaa atggatcagc tgcagaggac ggaagactta aagaaggtga 1920
tgaaatccta gatgtaaag gaataccaat aaagggttg acatttcaag aagccattca 1980
tacctttaag caaatccgga gtggattatt tgttttaacg gtacgcacaa agttggtgag 2040
ccccagctc acaccctgct cgacaccac acacatgagc agatccgcct cccgaactt 2100
caataccagt gggggagcct cggcgggagg ttccgatgaa ggcagttctt catccctggg 2160
tcggaagacc cctgggcca aggacaggat cgtcatggaa gtaacactca acaaagagcc 2220
aagagttgga ttaggcattg gtgcctgctg cttggctctg gaaaacagtc ctctggcat 2280
ctacattcac agccttgctc caggatcagt ggccaagatg gagagcaacc tgtcgcgggg 2340
atcaatcctg gaagtgaact ccgtcaacgt ccgccatgct gctttaagca aagtccacgc 2400
catcttgagt aaatgccctc caggaccctg tcgccttgct atcgccggc accctaattc 2460
aaaggtgaat caggtttccg agcaggaaat ggatgaagtc atagcacgca gcacttatca 2520
ggagagcaaa gagggcaatt cctctcctgg cttaggtact gtaatctcaa tcggatgttt 2580
tcttctcaa caggactccc ttatttctga atctgaactc tcccagtact ttgccacga 2640
tgtccctggc cccttgctcag acttcatggg ggccggttct gaggacgagg atcaccggg 2700
aagtggctgc agcacgtcgg aggagggcag cctgcctccc agcacctcca ctcacaagga 2760
gcctggaaaa cccagagcca acagcctcgt gactcttggg agccatcggg cttctgggct 2820

```

cttccacaag	caggtgacag	ttgccagaca	agccagtctc	cccggaagcc	cacaggccct	2880
ccgaaaccct	ctcctccgcc	agaggaaggt	aggtgctac	gatgccaacg	atgccagtga	2940
tgaggaagag	tttgacagag	aaggggactg	catttcactc	ccagggggccc	tcccgggtcc	3000
catcaggcct	ctgtcagagg	atgacccgag	gcggtgtctca	atttcctctt	ccaagggcat	3060
ggacgtccac	aaccaagagg	aacgaccccg	gaaaacactg	gtgagcaagg	ccatctcggc	3120
acctcttctt	ggtagctcag	tggacttaga	ggagagtatc	ccagagggga	tgggtgatgc	3180
tgcgtcctat	gcagccaacc	tcacggactc	tgcagaggcc	cccaagggga	gccctggaag	3240
ctggtggaag	aaggaactgt	caggatcaag	tagcgcaccc	aaattggaat	acacagtccg	3300
tacagacacc	cagagtccga	caaacactgg	gagccccagt	tccccccagc	aaaaaagtga	3360
aggcctgggc	tccaggcaca	gaccagtggc	cagggtaagc	ccccactgca	agagatccga	3420
ggctgaggcc	aagcccagtg	gctcacagac	agtgaacctg	actggcagag	ccaatgatcc	3480
atgcgatctg	gactcgagag	tccaggccac	ttctgtcaaa	gtgactgtcg	ctggctttca	3540
gccaggtgga	gctgtggaga	aggaatctct	gggaaagctg	accactggag	atgcttgtgt	3600
ctctaccagc	tgtgaactag	ccagtgtctt	gtcccatctg	gatgccagcc	acctcacaga	3660
gaacctgccc	aaagctgcat	cagagctggg	gcaacaaccc	atgactgaac	tggacagtc	3720
ctcggacctc	atctcttccc	caggaagaa	gggggcccgt	catcctgacc	ccagcaagac	3780
ctctgtagac	acagggaag	tcagtcggcc	agagaatccc	agccagcctg	catcgcccag	3840
ggtcgccaag	tgcaaggcca	ggtctccagt	caggctcccc	catgagggca	gccccctccc	3900
aggggagaaa	gcagcggctc	cccctgacta	cagcaagact	cgatcagcat	cggaaaccag	3960
cacaccccac	aataccagga	gggtggctgc	cctcagggga	gcgggacctg	gagcagaggg	4020
aatgacacca	gctggtgctg	tcctgccagg	agacccccctc	acatcccagg	agcagagaca	4080
gggagctcca	ggtaaccaca	gtaaggctct	ggaaatgaca	ggaatccatg	cacctgaaag	4140
ctcccaggag	ccttccctgc	tggagggagc	agattctgtg	tcctcaaggg	caccgcaggc	4200
cagcctctcc	atgttgccat	ccactgacaa	caccaaagaa	gcatgtggcc	atgtctcggg	4260
gcaactgtgc	ccggggggga	gtagagagag	ccctgtgacg	gacattgaca	gcttcatcaa	4320
ggagctggat	gcttctgcag	caaggtctcc	gtcttcccag	acgggggaca	gtggctctca	4380
ggagggcagt	gctcagggcc	accaccagc	cggggctgga	ggtgggagct	cctgccgtgc	4440
cgaaccagtc	ccggggggcc	agacctcctc	cccagggagg	gcctgggctg	ctggtgcccc	4500
cgctaccca	caatgggcct	cccagccttc	ggttttagat	tcaattaatc	ccgacaaaca	4560
ttttactgtg	aacaaaaact	ttctgagcaa	ctactctaga	aatttttagca	gttttcatga	4620
agacagcacc	tccttatcag	gcctgggtga	cagcacggag	ccgtctctgt	catccatgta	4680
tggcgatgct	gaggattctt	cttctgaccc	tgagtactc	actgaagccc	cacgagcttc	4740
tgccagggac	ggctgggtccc	ctcctcgttc	ccgtgtgtct	ttgcacaagg	aagatccttc	4800
ggagtcagaa	gaggaacaga	ttgagatttg	ttccacacgt	ggctgcccc	atccaccctc	4860
gagtctgct	catcttccca	cccaggctgc	catctgtcct	gcctcagcca	aagtctctgtc	4920
attaaaatac	agcactccga	gagagtcggt	ggccagtcct	cgtgagaagg	tcgctgctt	4980
gccaggctca	tacacttcag	gccagactc	ttcccagcca	tcactactct	tggagatgag	5040
ctctcaggag	catgaaactc	atgcggacat	aagcacttca	cagaaccaca	ggccctcgtg	5100
tgcagaagaa	accacagaag	tcaccagcgc	tagctcagcc	atggaaaaca	gtccgctgtc	5160
taaagtagcc	aggcattttc	acagtccgcc	catcattctc	agctcccca	acatggtaaa	5220
tggcttgga	catgacctgc	tagatgacga	aacctgaat	caatacgaaa	caagcattaa	5280
tgcagctgcc	agtctgtcct	ccttcagtgt	ggatgtccct	aagaatggag	aatctgtttt	5340
ggaaaacctc	cacatctctg	aaagtcaaga	cctggatgac	ttgctacaga	aacaaaaaat	5400
gatcgctagg	aggcccatca	tggcctgggt	taaagaaata	aataaacata	accaaggcac	5460
acatttgagg	agcaaaaccg	agaaggaaca	acctctaata	cctgccagaa	gtcccgaactc	5520
caagattcag	atggtgagtt	caagccaaaa	aaagggcggt	actgtgcctc	atagccctcc	5580
tcagccgaaa	acaaacctgg	aaaataagga	cctgtctaa	aagagtccgg	cagaaatgct	5640
tctgactaat	ggtcagaagg	caaagtgtgg	tccgaagctg	aagaggctca	gcctcaaggg	5700
caaggccaaa	gtcaactctg	aggccccctgc	tgcgaatgct	gtgaaggctg	gggggacgga	5760
ccacaggaaa	cccttgatct	caccccagac	ctcccacaaa	acactttcta	aggcagtgtc	5820
acagcggctc	catgtagccg	accacgagga	ccctgacaga	aacaccacag	ctgccccag	5880
gtccccccag	tgtgtgctgg	aaagcaagcc	acctcttgcc	acctctgggc	cactgaaacc	5940
ctcagtgtct	gacacgagca	tcaggacatt	tgtctcgccc	ctgacctctc	ccaagcctgt	6000
tcctgagcaa	ggcatgtgga	gcaggttcca	catggctgtc	ctctctgaac	ccgacagagg	6060
ttgcccaacc	accctaaat	ctcctaagtg	tagagcagag	ggcagggcgc	cccgtgctga	6120
ctccgggccc	gtgagtccgg	cagcgtctag	gaacggcatg	tccgtggcag	ggaacagaca	6180
gagtgaagcc	cgctggcca	gccatgtggc	agcagacaca	gccaaccca	ggccgactgg	6240

cgaaaaagga ggcaacataa tggccagcga tgcctcgaa agaacaacc agctgaaaat 6300
 cgtggagatt tctgctgaag cagtgtcaga gactgtatgt ggtaacaagc cagctgaaag 6360
 cgacagacgg ggaggggtgct tggcccaggg caactgtcag gagaagagtg aaatcaggct 6420
 ctatcgccag gtcgcagaat catccacaag tcatccatcc tcaatcccat ctcatgcctc 6480
 ccaggcagag caggaaatgt cacgatcatt cagcatggca aaactggcgt cctcctcctc 6540
 ctcccttcaa acagccatta gaaaggcaga atactcccag ggaaaatcaa gcctgatgtc 6600
 agactcccga ggggtgcca gaaacagcat tccagggggc ccctcggggg aggaccatct 6660
 ctacttcacc ccaaggccag cgaccaggac ctactccatg ccagcccagt tctcaagcca 6720
 ttttgagcgg gagggtcacc cccacacag cctgggtcgc tctcgggaca gccagggtccc 6780
 tgtgacaagc agtgttgctc ccgaggcaaa ggcattccaga ggtggtcttc ccagcctggc 6840
 taatggacag ggcatatata gtgtaaagcc gctgctggac acatcgagga atcttccagc 6900
 cacagatgaa ggggatata tttcagtcga ggagacgagc tgcctagtca cagacaaaat 6960
 caaagtccac agacgacact actgctatga gcagaactgg ccccatgaat ctacctcatt 7020
 tttctctgtg aagcagcggg tcaagtcttt tgagaacctg gccaatgctg accggcctgt 7080
 agccaagtcc ggggcttccc catttttgtc ggtgagctcc aagcctcca ttgggaggcg 7140
 gtcttccggc agcattgttt ccgggagcct gggccacca ggtgacgcag cagcaaggtt 7200
 gttgagacgc agcttgagtt cctgcagcga aaaccaaagc gaagccggca ccctcctgcc 7260
 ccagatggcc aagtctccct caatcatgac actgaccatc tctcggcaga acccaccaga 7320
 gaccagtagc aagggtctctg attcggaaact aaagaaatca cttggtcctt tgggaattcc 7380
 caccacaacg atgacctggt cttctcctgt taagaggaac aagtcctcgg tacgccacac 7440
 gcagccctcg cccgtgtccc gctccaagct ccaggagctg agagccttga gcatgcctga 7500
 ccttgacaag ctctgcagcg aggattactc agcagggccg agcgcctgct tcttcaaaac 7560
 tgagctggag atcaccccca ggaggtcacc tggccctcct gctggaggcg tttcgtgtcc 7620
 cgagaagggc ggggaacaggg cctgtccagg aggaagtggc cctaaaacca gtgctgctga 7680
 gacaccaggt tcagccagtg atacgggtga agctgcccag gatctgcctt ttagaagaag 7740
 ctggctcagtt aatttgatc aacttctagt ctcagcgggg gaccagcaa gattacagtc 7800
 tgttttatcg tcagtgggat cgaaatctac catcctaact ctcattcagg aagcgaagc 7860
 acaatcagag aatgaagaag atgtttgctt catagtcttg aatagaaaag aaggctcagg 7920
 tctgggattc agtgtggcag gagggacaga tgtggagcca aaatcaatca cggccacag 7980
 ggtgttttct cagggggcgg cttctcagga agggactatg aaccgagggg atttccttct 8040
 gtcagtcaac ggcgcctcac tggctggcct agcccacggg aatgtcctga aggttctgca 8100
 ccaggcacag ctgcacaaag atgccctcgt ggtcatcaag aaagggatgg atcagcccag 8160
 gccctctgcc cggcaggagc ctcccacagc caatgggaag ggtttgctgt ccagaaagac 8220
 catccccctg gagcctggca ttgggagaag tgtggctgta cacgatgctc tgtgtgttga 8280
 agtgcgtgaag acctcggctg ggctgggact gagtctggat gggggaaaat catcgggtgac 8340
 gggagatggg cccttggtca ttaaaagagt gtacaaaggt ggtgcggctg aacaagctgg 8400
 aataatagaa gctggagatg aaattcttgc tattaatggg aaacctctgg ttgggctcat 8460
 gcactttgat gcctggaata ttatgaagtc tgtcccagaa ggacctgtgc agttattaat 8520
 tagaaagcat aggaattctt catgaatttt aacaagaatc attttctcag ttctcttctt 8580
 tctttagcaa atcagagtga cttctttaa ccacaggttg ttgaaatggc caacactggg 8640

<210> 25

<211> 2814

<212> PRT

<213> Homo sapiens

<400> 25

Met	Pro	Ile	Thr	Gln	Asp	Asn	Ala	Val	Leu	His	Leu	Pro	Leu	Leu	Tyr
1				5					10					15	
Gln	Trp	Leu	Gln	Asn	Ser	Leu	Gln	Glu	Gly	Gly	Asp	Gly	Pro	Glu	Gln
			20					25					30		
Arg	Leu	Cys	Gln	Ala	Ala	Ile	Gln	Lys	Leu	Gln	Glu	Tyr	Ile	Gln	Leu
		35					40					45			

Asn Phe Ala Val Asp Glu Ser Thr Val Pro Pro Asp His Ser Pro Pro
 50 55 60
 Glu Met Glu Ile Cys Thr Val Tyr Leu Thr Lys Glu Leu Gly Asp Thr
 65 70 75 80
 Glu Thr Val Gly Leu Ser Phe Gly Asn Ile Pro Val Phe Gly Asp Tyr
 85 90 95
 Gly Glu Lys Arg Arg Gly Gly Lys Lys Arg Lys Thr His Gln Gly Pro
 100 105 110
 Val Leu Asp Val Gly Cys Ile Trp Val Thr Glu Leu Arg Lys Asn Ser
 115 120 125
 Pro Ala Gly Lys Ser Gly Lys Val Arg Leu Arg Asp Glu Ile Leu Ser
 130 135 140
 Leu Asn Gly Gln Leu Met Val Gly Val Asp Val Ser Gly Ala Ser Tyr
 145 150 155 160
 Leu Ala Glu Gln Cys Trp Asn Gly Gly Phe Ile Tyr Leu Ile Met Leu
 165 170 175
 Arg Arg Phe Lys His Lys Ala His Ser Thr Tyr Asn Gly Asn Ser Ser
 180 185 190
 Asn Ser Ser Glu Pro Gly Glu Thr Pro Thr Leu Glu Leu Gly Asp Arg
 195 200 205
 Thr Ala Lys Lys Gly Lys Arg Thr Arg Lys Phe Gly Val Ile Ser Arg
 210 215 220
 Pro Pro Ala Asn Lys Ala Pro Glu Glu Ser Lys Gly Ser Ala Gly Cys
 225 230 235 240
 Glu Val Ser Ser Asp Pro Ser Thr Glu Leu Glu Asn Gly Leu Asp Pro
 245 250 255
 Glu Leu Gly Asn Gly His Val Phe Gln Leu Glu Asn Gly Pro Asp Ser
 260 265 270
 Leu Lys Glu Val Ala Gly Pro His Leu Glu Arg Ser Glu Val Asp Arg
 275 280 285
 Gly Thr Glu His Arg Ile Pro Lys Thr Asp Ala Pro Leu Thr Thr Ser
 290 295 300
 Asn Asp Lys Arg Arg Phe Ser Lys Gly Gly Lys Thr Asp Phe Gln Ser
 305 310 315 320
 Ser Asp Cys Leu Ala Arg Ser Lys Glu Glu Val Gly Arg Ile Trp Lys
 325 330 335
 Met Glu Leu Leu Lys Glu Ser Asp Gly Leu Gly Ile Gln Val Ser Gly
 340 345 350

Gly Arg Gly Ser Lys Arg Ser Pro His Ala Ile Val Val Thr Gln Val
 355 360 365
 Lys Glu Gly Gly Ala Ala His Arg Leu Arg Asp Gly Arg Leu Ser Leu
 370 375 380
 Gly Asp Glu Leu Leu Val Ile Asn Gly His Leu Leu Val Gly Leu Ser
 385 390 395 400
 His Glu Glu Ala Val Ala Ile Leu Arg Ser Ala Thr Gly Met Val Gln
 405 410 415
 Leu Val Val Ala Ser Lys Val Gly Val Leu Ser Ala Phe Gln Met Pro
 420 425 430
 Gly Thr Asp Glu Pro Gln Asp Val Cys Gly Ala Glu Glu Ser Lys Gly
 435 440 445
 Asn Leu Glu Ser Pro Lys Gln Gly Ser Asn Lys Ile Lys Leu Lys Ser
 450 455 460
 Arg Leu Ser Gly Arg Trp Gly Leu Tyr Leu Met Gln Pro Val Gly Gly
 465 470 475 480
 Val His Arg Leu Glu Ser Val Glu Glu Tyr Asn Glu Leu Met Val Arg
 485 490 495
 Asn Gly Asp Pro Arg Ile Arg Met Leu Glu Val Ser Arg Asp Gly Arg
 500 505 510
 Lys His Ser Leu Pro Gln Leu Leu Asp Ser Ser Ser Ala Ser Gln Glu
 515 520 525
 Tyr His Ile Val Lys Lys Ser Thr Arg Ser Leu Ser Thr Thr Gln Val
 530 535 540
 Glu Ser Pro Trp Arg Leu Ile Arg Pro Ser Val Ile Ser Ile Ile Gly
 545 550 555 560
 Leu Tyr Lys Glu Lys Gly Lys Gly Leu Gly Phe Ser Ile Ala Gly Gly
 565 570 575
 Arg Asp Cys Ile Arg Gly Gln Met Gly Ile Phe Val Lys Thr Ile Phe
 580 585 590
 Pro Asn Gly Ser Ala Ala Glu Asp Gly Arg Leu Lys Glu Gly Asp Glu
 595 600 605
 Ile Leu Asp Val Asn Gly Ile Pro Ile Lys Gly Leu Thr Phe Gln Glu
 610 615 620
 Ala Ile His Thr Phe Lys Gln Ile Arg Ser Gly Leu Phe Val Leu Thr
 625 630 635 640
 Val Arg Thr Lys Leu Val Ser Pro Ser Leu Thr Pro Cys Ser Thr Pro
 645 650 655

Thr His Met Ser Arg Ser Ala Ser Pro Asn Phe Asn Thr Ser Gly Gly
 660 665 670
 Ala Ser Ala Gly Gly Ser Asp Glu Gly Ser Ser Ser Ser Leu Gly Arg
 675 680 685
 Lys Thr Pro Gly Pro Lys Asp Arg Ile Val Met Glu Val Thr Leu Asn
 690 695 700
 Lys Glu Pro Arg Val Gly Leu Gly Ile Gly Ala Cys Cys Leu Ala Leu
 705 710 715 720
 Glu Asn Ser Pro Pro Gly Ile Tyr Ile His Ser Leu Ala Pro Gly Ser
 725 730 735
 Val Ala Lys Met Glu Ser Asn Leu Ser Arg Gly Ser Ile Leu Glu Val
 740 745 750
 Asn Ser Val Asn Val Arg His Ala Ala Leu Ser Lys Val His Ala Ile
 755 760 765
 Leu Ser Lys Cys Pro Pro Gly Pro Val Arg Leu Val Ile Gly Arg His
 770 775 780
 Pro Asn Pro Lys Val Asn Gln Val Ser Glu Gln Glu Met Asp Glu Val
 785 790 795 800
 Ile Ala Arg Ser Thr Tyr Gln Glu Ser Lys Glu Ala Asn Ser Ser Pro
 805 810 815
 Gly Leu Gly Thr Val Ile Ser Ile Gly Cys Phe Leu Leu Gln Gln Asp
 820 825 830
 Ser Leu Ile Ser Glu Ser Glu Leu Ser Gln Tyr Phe Ala His Asp Val
 835 840 845
 Pro Gly Pro Leu Ser Asp Phe Met Val Ala Gly Ser Glu Asp Glu Asp
 850 855 860
 His Pro Gly Ser Gly Cys Ser Thr Ser Glu Glu Gly Ser Leu Pro Pro
 865 870 875 880
 Ser Thr Ser Thr His Lys Glu Pro Gly Lys Pro Arg Ala Asn Ser Leu
 885 890 895
 Val Thr Leu Gly Ser His Arg Ala Ser Gly Leu Phe His Lys Gln Val
 900 905 910
 Thr Val Ala Arg Gln Ala Ser Leu Pro Gly Ser Pro Gln Ala Leu Arg
 915 920 925
 Asn Pro Leu Leu Arg Gln Arg Lys Val Gly Cys Tyr Asp Ala Asn Asp
 930 935 940
 Ala Ser Asp Glu Glu Glu Phe Asp Arg Glu Gly Asp Cys Ile Ser Leu
 945 950 955 960

Pro Gly Ala Leu Pro Gly Pro Ile Arg Pro Leu Ser Glu Asp Asp Pro
 965 970 975

Arg Arg Val Ser Ile Ser Ser Ser Lys Gly Met Asp Val His Asn Gln
 980 985 990

Glu Glu Arg Pro Arg Lys Thr Leu Val Ser Lys Ala Ile Ser Ala Pro
 995 1000 1005

Leu Leu Gly Ser Ser Val Asp Leu Glu Glu Ser Ile Pro Glu Gly Met
 1010 1015 1020

Val Asp Ala Ala Ser Tyr Ala Ala Asn Leu Thr Asp Ser Ala Glu Ala
 1025 1030 1035 1040

Pro Lys Gly Ser Pro Gly Ser Trp Trp Lys Lys Glu Leu Ser Gly Ser
 1045 1050 1055

Ser Ser Ala Pro Lys Leu Glu Tyr Thr Val Arg Thr Asp Thr Gln Ser
 1060 1065 1070

Pro Thr Asn Thr Gly Ser Pro Ser Ser Pro Gln Gln Lys Ser Glu Gly
 1075 1080 1085

Leu Gly Ser Arg His Arg Pro Val Ala Arg Val Ser Pro His Cys Lys
 1090 1095 1100

Arg Ser Glu Ala Glu Ala Lys Pro Ser Gly Ser Gln Thr Val Asn Leu
 1105 1110 1115 1120

Thr Gly Arg Ala Asn Asp Pro Cys Asp Leu Asp Ser Arg Val Gln Ala
 1125 1130 1135

Thr Ser Val Lys Val Thr Val Ala Gly Phe Gln Pro Gly Gly Ala Val
 1140 1145 1150

Glu Lys Glu Ser Leu Gly Lys Leu Thr Thr Gly Asp Ala Cys Val Ser
 1155 1160 1165

Thr Ser Cys Glu Leu Ala Ser Ala Leu Ser His Leu Asp Ala Ser His
 1170 1175 1180

Leu Thr Glu Asn Leu Pro Lys Ala Ala Ser Glu Leu Gly Gln Gln Pro
 1185 1190 1195 1200

Met Thr Glu Leu Asp Ser Ser Ser Asp Leu Ile Ser Ser Pro Gly Lys
 1205 1210 1215

Lys Gly Ala Ala His Pro Asp Pro Ser Lys Thr Ser Val Asp Thr Gly
 1220 1225 1230

Lys Val Ser Arg Pro Glu Asn Pro Ser Gln Pro Ala Ser Pro Arg Val
 1235 1240 1245

Ala Lys Cys Lys Ala Arg Ser Pro Val Arg Leu Pro His Glu Gly Ser
 1250 1255 1260

Pro Ser Pro Gly Glu Lys Ala Ala Ala Pro Pro Asp Tyr Ser Lys Thr
 1265 1270 1275 1280
 Arg Ser Ala Ser Glu Thr Ser Thr Pro His Asn Thr Arg Arg Val Ala
 1285 1290 1295
 Ala Leu Arg Gly Ala Gly Pro Gly Ala Glu Gly Met Thr Pro Ala Gly
 1300 1305 1310
 Ala Val Leu Pro Gly Asp Pro Leu Thr Ser Gln Glu Gln Arg Gln Gly
 1315 1320 1325
 Ala Pro Gly Asn His Ser Lys Ala Leu Glu Met Thr Gly Ile His Ala
 1330 1335 1340
 Pro Glu Ser Ser Gln Glu Pro Ser Leu Leu Glu Gly Ala Asp Ser Val
 1345 1350 1355 1360
 Ser Ser Arg Ala Pro Gln Ala Ser Leu Ser Met Leu Pro Ser Thr Asp
 1365 1370 1375
 Asn Thr Lys Glu Ala Cys Gly His Val Ser Gly His Cys Cys Pro Gly
 1380 1385 1390
 Gly Ser Arg Glu Ser Pro Val Thr Asp Ile Asp Ser Phe Ile Lys Glu
 1395 1400 1405
 Leu Asp Ala Ser Ala Ala Arg Ser Pro Ser Ser Gln Thr Gly Asp Ser
 1410 1415 1420
 Gly Ser Gln Glu Gly Ser Ala Gln Gly His Pro Pro Ala Gly Ala Gly
 1425 1430 1435 1440
 Gly Gly Ser Ser Cys Arg Ala Glu Pro Val Pro Gly Gly Gln Thr Ser
 1445 1450 1455
 Ser Pro Arg Arg Ala Trp Ala Ala Gly Ala Pro Ala Tyr Pro Gln Trp
 1460 1465 1470
 Ala Ser Gln Pro Ser Val Leu Asp Ser Ile Asn Pro Asp Lys His Phe
 1475 1480 1485
 Thr Val Asn Lys Asn Phe Leu Ser Asn Tyr Ser Arg Asn Phe Ser Ser
 1490 1495 1500
 Phe His Glu Asp Ser Thr Ser Leu Ser Gly Leu Gly Asp Ser Thr Glu
 1505 1510 1515 1520
 Pro Ser Leu Ser Ser Met Tyr Gly Asp Ala Glu Asp Ser Ser Ser Asp
 1525 1530 1535
 Pro Glu Ser Leu Thr Glu Ala Pro Arg Ala Ser Ala Arg Asp Gly Trp
 1540 1545 1550
 Ser Pro Pro Arg Ser Arg Val Ser Leu His Lys Glu Asp Pro Ser Glu
 1555 1560 1565

Ser Glu Glu Glu Gln Ile Glu Ile Cys Ser Thr Arg Gly Cys Pro Asn
 1570 1575 1580

Pro Pro Ser Ser Pro Ala His Leu Pro Thr Gln Ala Ala Ile Cys Pro
 1585 1590 1595 1600

Ala Ser Ala Lys Val Leu Ser Leu Lys Tyr Ser Thr Pro Arg Glu Ser
 1605 1610 1615

Val Ala Ser Pro Arg Glu Lys Val Ala Cys Leu Pro Gly Ser Tyr Thr
 1620 1625 1630

Ser Gly Pro Asp Ser Ser Gln Pro Ser Ser Leu Leu Glu Met Ser Ser
 1635 1640 1645

Gln Glu His Glu Thr His Ala Asp Ile Ser Thr Ser Gln Asn His Arg
 1650 1655 1660

Pro Ser Cys Ala Glu Glu Thr Thr Glu Val Thr Ser Ala Ser Ser Ala
 1665 1670 1675 1680

Met Glu Asn Ser Pro Leu Ser Lys Val Ala Arg His Phe His Ser Pro
 1685 1690 1695

Pro Ile Ile Leu Ser Ser Pro Asn Met Val Asn Gly Leu Glu His Asp
 1700 1705 1710

Leu Leu Asp Asp Glu Thr Leu Asn Gln Tyr Glu Thr Ser Ile Asn Ala
 1715 1720 1725

Ala Ala Ser Leu Ser Ser Phe Ser Val Asp Val Pro Lys Asn Gly Glu
 1730 1735 1740

Ser Val Leu Glu Asn Leu His Ile Ser Glu Ser Gln Asp Leu Asp Asp
 1745 1750 1755 1760

Leu Leu Gln Lys Pro Lys Met Ile Ala Arg Arg Pro Ile Met Ala Trp
 1765 1770 1775

Phe Lys Glu Ile Asn Lys His Asn Gln Gly Thr His Leu Arg Ser Lys
 1780 1785 1790

Thr Glu Lys Glu Gln Pro Leu Met Pro Ala Arg Ser Pro Asp Ser Lys
 1795 1800 1805

Ile Gln Met Val Ser Ser Ser Gln Lys Lys Gly Val Thr Val Pro His
 1810 1815 1820

Ser Pro Pro Gln Pro Lys Thr Asn Leu Glu Asn Lys Asp Leu Ser Lys
 1825 1830 1835 1840

Lys Ser Pro Ala Glu Met Leu Leu Thr Asn Gly Gln Lys Ala Lys Cys
 1845 1850 1855

Gly Pro Lys Leu Lys Arg Leu Ser Leu Lys Gly Lys Ala Lys Val Asn
 1860 1865 1870

Ser Glu Ala Pro Ala Ala Asn Ala Val Lys Ala Gly Gly Thr Asp His
 1875 1880 1885

Arg Lys Pro Leu Ile Ser Pro Gln Thr Ser His Lys Thr Leu Ser Lys
 1890 1895 1900

Ala Val Ser Gln Arg Leu His Val Ala Asp His Glu Asp Pro Asp Arg
 1905 1910 1915 1920

Asn Thr Thr Ala Ala Pro Arg Ser Pro Gln Cys Val Leu Glu Ser Lys
 1925 1930 1935

Pro Pro Leu Ala Thr Ser Gly Pro Leu Lys Pro Ser Val Ser Asp Thr
 1940 1945 1950

Ser Ile Arg Thr Phe Val Ser Pro Leu Thr Ser Pro Lys Pro Val Pro
 1955 1960 1965

Glu Gln Gly Met Trp Ser Arg Phe His Met Ala Val Leu Ser Glu Pro
 1970 1975 1980

Asp Arg Gly Cys Pro Thr Thr Pro Lys Ser Pro Lys Cys Arg Ala Glu
 1985 1990 1995 2000

Gly Arg Ala Pro Arg Ala Asp Ser Gly Pro Val Ser Pro Ala Ala Ser
 2005 2010 2015

Arg Asn Gly Met Ser Val Ala Gly Asn Arg Gln Ser Glu Pro Arg Leu
 2020 2025 2030

Ala Ser His Val Ala Ala Asp Thr Ala Gln Pro Arg Pro Thr Gly Glu
 2035 2040 2045

Lys Gly Gly Asn Ile Met Ala Ser Asp Arg Leu Glu Arg Thr Asn Gln
 2050 2055 2060

Leu Lys Ile Val Glu Ile Ser Ala Glu Ala Val Ser Glu Thr Val Cys
 2065 2070 2075 2080

Gly Asn Lys Pro Ala Glu Ser Asp Arg Arg Gly Gly Cys Leu Ala Gln
 2085 2090 2095

Gly Asn Cys Gln Glu Lys Ser Glu Ile Arg Leu Tyr Arg Gln Val Ala
 2100 2105 2110

Glu Ser Ser Thr Ser His Pro Ser Ser Leu Pro Ser His Ala Ser Gln
 2115 2120 2125

Ala Glu Gln Glu Met Ser Arg Ser Phe Ser Met Ala Lys Leu Ala Ser
 2130 2135 2140

Ser Ser Ser Ser Leu Gln Thr Ala Ile Arg Lys Ala Glu Tyr Ser Gln
 2145 2150 2155 2160

Gly Lys Ser Ser Leu Met Ser Asp Ser Arg Gly Val Pro Arg Asn Ser
 2165 2170 2175

Ile Pro Gly Gly Pro Ser Gly Glu Asp His Leu Tyr Phe Thr Pro Arg
 2180 2185 2190
 Pro Ala Thr Arg Thr Tyr Ser Met Pro Ala Gln Phe Ser Ser His Phe
 2195 2200 2205
 Gly Arg Glu Gly His Pro Pro His Ser Leu Gly Arg Ser Arg Asp Ser
 2210 2215 2220
 Gln Val Pro Val Thr Ser Ser Val Val Pro Glu Ala Lys Ala Ser Arg
 2225 2230 2235 2240
 Gly Gly Leu Pro Ser Leu Ala Asn Gly Gln Gly Ile Tyr Ser Val Lys
 2245 2250 2255
 Pro Leu Leu Asp Thr Ser Arg Asn Leu Pro Ala Thr Asp Glu Gly Asp
 2260 2265 2270
 Ile Ile Ser Val Gln Glu Thr Ser Cys Leu Val Thr Asp Lys Ile Lys
 2275 2280 2285
 Val Thr Arg Arg His Tyr Cys Tyr Glu Gln Asn Trp Pro His Glu Ser
 2290 2295 2300
 Thr Ser Phe Phe Ser Val Lys Gln Arg Ile Lys Ser Phe Glu Asn Leu
 2305 2310 2315 2320
 Ala Asn Ala Asp Arg Pro Val Ala Lys Ser Gly Ala Ser Pro Phe Leu
 2325 2330 2335
 Ser Val Ser Ser Lys Pro Pro Ile Gly Arg Arg Ser Ser Gly Ser Ile
 2340 2345 2350
 Val Ser Gly Ser Leu Gly His Pro Gly Asp Ala Ala Ala Arg Leu Leu
 2355 2360 2365
 Arg Arg Ser Leu Ser Ser Cys Ser Glu Asn Gln Ser Glu Ala Gly Thr
 2370 2375 2380
 Leu Leu Pro Gln Met Ala Lys Ser Pro Ser Ile Met Thr Leu Thr Ile
 2385 2390 2395 2400
 Ser Arg Gln Asn Pro Pro Glu Thr Ser Ser Lys Gly Ser Asp Ser Glu
 2405 2410 2415
 Leu Lys Lys Ser Leu Gly Pro Leu Gly Ile Pro Thr Pro Thr Met Thr
 2420 2425 2430
 Leu Ala Ser Pro Val Lys Arg Asn Lys Ser Ser Val Arg His Thr Gln
 2435 2440 2445
 Pro Ser Pro Val Ser Arg Ser Lys Leu Gln Glu Leu Arg Ala Leu Ser
 2450 2455 2460
 Met Pro Asp Leu Asp Lys Leu Cys Ser Glu Asp Tyr Ser Ala Gly Pro
 2465 2470 2475 2480

Ser Ala Val Leu Phe Lys Thr Glu Leu Glu Ile Thr Pro Arg Arg Ser
 2485 2490 2495
 Pro Gly Pro Pro Ala Gly Gly Val Ser Cys Pro Glu Lys Gly Gly Asn
 2500 2505 2510
 Arg Ala Cys Pro Gly Gly Ser Gly Pro Lys Thr Ser Ala Ala Glu Thr
 2515 2520 2525
 Pro Ser Ser Ala Ser Asp Thr Gly Glu Ala Ala Gln Asp Leu Pro Phe
 2530 2535 2540
 Arg Arg Ser Trp Ser Val Asn Leu Asp Gln Leu Leu Val Ser Ala Gly
 2545 2550 2555 2560
 Asp Gln Gln Arg Leu Gln Ser Val Leu Ser Ser Val Gly Ser Lys Ser
 2565 2570 2575
 Thr Ile Leu Thr Leu Ile Gln Glu Ala Lys Ala Gln Ser Glu Asn Glu
 2580 2585 2590
 Glu Asp Val Cys Phe Ile Val Leu Asn Arg Lys Glu Gly Ser Gly Leu
 2595 2600 2605
 Gly Phe Ser Val Ala Gly Gly Thr Asp Val Glu Pro Lys Ser Ile Thr
 2610 2615 2620
 Val His Arg Val Phe Ser Gln Gly Ala Ala Ser Gln Glu Gly Thr Met
 2625 2630 2635 2640
 Asn Arg Gly Asp Phe Leu Leu Ser Val Asn Gly Ala Ser Leu Ala Gly
 2645 2650 2655
 Leu Ala His Gly Asn Val Leu Lys Val Leu His Gln Ala Gln Leu His
 2660 2665 2670
 Lys Asp Ala Leu Val Val Ile Lys Lys Gly Met Asp Gln Pro Arg Pro
 2675 2680 2685
 Ser Ala Arg Gln Glu Pro Pro Thr Ala Asn Gly Lys Gly Leu Leu Ser
 2690 2695 2700
 Arg Lys Thr Ile Pro Leu Glu Pro Gly Ile Gly Arg Ser Val Ala Val
 2705 2710 2715 2720
 His Asp Ala Leu Cys Val Glu Val Leu Lys Thr Ser Ala Gly Leu Gly
 2725 2730 2735
 Leu Ser Leu Asp Gly Gly Lys Ser Ser Val Thr Gly Asp Gly Pro Leu
 2740 2745 2750
 Val Ile Lys Arg Val Tyr Lys Gly Gly Ala Ala Glu Gln Ala Gly Ile
 2755 2760 2765
 Ile Glu Ala Gly Asp Glu Ile Leu Ala Ile Asn Gly Lys Pro Leu Val
 2770 2775 2780

Gly Leu Met His Phe Asp Ala Trp Asn Ile Met Lys Ser Val Pro Glu
 2785 2790 2795 2800

Gly Pro Val Gln Leu Leu Ile Arg Lys His Arg Asn Ser Ser
 2805 2810

<210> 26
 <211> 8640
 <212> DNA
 <213> Homo sapiens

<400> 26

```

agctgatgat ggccagggac cccaggggac gtgggggcct gtgggggtctg gccccagga 60
gcaagacctc tgatgatgct ggtgtctggg agtgagcacc atgcccacatca cccaggacaa 120
tgccgtgctg cacctgcccc tcctctacca gtggctgcag aacagcctgc aggaagggtg 180
ggatggggccg gagcagcggc tctgccaggc ggccatccag aagctgcagg agtacatcca 240
gctgaacttt gctgtggatg agagtacggt cccacctgat cacagcccc ccgaaatgga 300
gatctgtact gtgtacctca ccaaggagct gggggacaca gagactgtgg gcctgagttt 360
tggaacatc cctgttttcg gggactatgg tgaagagcgc agggggggca agaagaggaa 420
aaccaccag ggtcctgtgc tggatgtggg ctgcatctgg gtgacagagc tgaggaaaga 480
cagcccagca ggaagagtg ggaagggtccg actgcgggat gagatcctct cactgaatgg 540
gcagctgatg gttggagttg atgtcagtgg ggccagttac ctggctgagc agtgctggaa 600
tgccggttt atctacctga tcatgtgcg tcgctttaag cacaagccc actccactta 660
taatggcaac agtagcaaca gctctgaacc aggagaaaca cctaccttgg agctgggtga 720
ccgaactgcg aaaaagggga aacgaaccag aaagtttggg gtcactctca ggcctcctgc 780
caacaaggcc cctgaagaat ccaagggcag cgctggctgt gaggtgtcca gtgaccccag 840
cactgagctg gagaacggcc tggaccctga acttggaac ggccatgtct ttcagctaga 900
aaatggccca gattctctca aggaggtggc tggaccccat ctagagaggt cagaagtgga 960
cagagggaca gagcatagaa ttccaaagac agatgtcct ctgaccacaa gcaatgaca 1020
acgccgttc tcaaaagggtg ggaagacgga cttccaatcg agtgactgcc tggcacgggc 1080
caaggaggaa gttggccgaa tatggaagat ggagctgctc aaagaatcgg atgggctggg 1140
aattcaggtt agtggaggcc gaggatcaaa gcgctcacct cacgctatcg ttgtcactca 1200
agtgaaggaa ggaggtgccg ctcacaggct cagggatggc aggtgtcct taggagatga 1260
gctgctggta atcaatggc atttactggt cgggctctcc cacgaggaag cagtggccat 1320
tcttcgctcc gccacgggaa tgggtgcagc tgtggtggcc agcaaggtag gtgtgctttc 1380
tgcatctcag atgcctggga cagatgaacc ccaagatgtg tgcggtgctg aggaatccaa 1440
ggggaacttg gaaagtccca aacagggcag caataaaatc aagctcaaga gtcgcctttc 1500
aggtaggtgg gggctctacc tgatgcagcc tgtcgggggt gtacaccgcc ttgagtcagt 1560
tgaagaatat aacgagctga tgggtcggaa tggggacccc cggatccgga tgttgagggt 1620
ctcccagat ggccggaaac actccctccc ctccttaagc acgactcagg tggaatctcc 1740
ggaataccac attgtgaaga agtctaccgc ctccttaagc acgactcagg tggaatctcc 1740
tcggaggctc attcggccat ccgtcatctc gatcattggg ttgtacaaag aaaaaggcaa 1800
gggccttggc tttagtattg ctggaggtcg agactgcatt cgtggacaga tggggatttt 1860
tgtcaagacc atcttcccaa atggatcagc tgcagaggac ggaagactta aagaagggga 1920
tgaaatccta gatgtaaag gaataaccaat aaagggttg acatttcaag aagccattca 1980
tacctttaag caaatccgga gtggattatt tgttttaacg gtacgcacaa agttggtgag 2040
ccccagcctc acaccctgct cgacaccac acacatgagc agatccgcct ccccgaaactt 2100
caataccagt gggggagcct cggcgggagg ttccgatgaa ggcagttctt catccctggg 2160
tcggaagacc cctgggcca aggacaggat cgtcatggaa gtaacactca acaaagagcc 2220
aagagttgga ttaggcattg gtgcctgctg cttggctctg gaaaacagtc ctctggcat 2280
ctacattcac agccttgctc caggatcagt ggccaagatg gagagcaacc tgagccgagg 2340
ggatcaaact ctggaagtga actccgtcaa cgtccgccat gctgctttaa gcaaagtcca 2400
cgcatcttg agtaaagcc ctccaggacc cgttcgcctt gtcacggcc ggcaccctaa 2460
tccaaagggt tccgagcagg aaatggatga agtcatagca cgcagcactt atcaggagag 2520
caaagaggcc aattcctctc ctggcttagg taccctcttg aagagtcctt ctcttgcaaa 2580
aaaggactcc cttatttctg aatctgaact ctcccagtag tttgccacg atgtccctgg 2640

```

ccccttgtca	gacttcatgg	tggtcggttc	tgaggacgag	gatcaccg	gaagtggctg	2700
cagcacgtcg	gaggagggca	gcctgcctcc	cagcacctcc	actcacaagg	agcctggaaa	2760
accagagacc	aacagcctcg	tgactcttgg	gagccatcgg	gcttctgggc	tcttccacaa	2820
gcaggtgaca	gttgccagac	aagccagtct	ccccggaagc	ccacaggccc	tccgaaaccc	2880
tctcctccgc	cagaggaagg	taggctgcta	cgatgccaac	gatgccagt	atgaggaaga	2940
gtttgacaga	gaaggggact	gcatttctact	cccagggggc	ctcccgggtc	ccatcaggcc	3000
tctgtcagag	gatgaccgca	ggcgtgtctc	aatttctct	tccaagggca	tggacgtcca	3060
caaccaagag	gaacgacccc	ggaaaacact	ggtgagcaag	gccatctcgg	cacctcttct	3120
tggtagctca	gtggacttag	aggagagtat	cccagagggc	atggtggatg	ctgcgtccta	3180
tgcagccaac	ctcacggact	ctgcagaggc	ccccaaaggg	agccctggaa	gctggtggaa	3240
gaaggaactg	tcaggatcaa	gtagcgcacc	caaattggaa	tacacagtcc	gtacagacac	3300
ccagagtcgg	acaaacactg	ggagccccag	ttccccccag	caaaaaagt	aaggcctggg	3360
ctccaggcac	agaccagtgg	ccagggtaa	ccccactgc	aagagatccg	aggctgaggc	3420
caagcccagt	ggctcacaga	cagtgaacct	gactggcaga	gccaatgatc	catgcgatct	3480
ggactcgaga	gtccaggcca	cttctgtcaa	agtactgtc	gctggcttct	agccagggtg	3540
agctgtggag	aaggaatctc	tgggaaagct	gaccactgga	gatgcttggt	tctctaccag	3600
ctgtgaacta	gccagtgtc	tgtcccatct	ggatgccagc	cacctcacag	agaacctgcc	3660
caaagctgca	tcagagctgg	ggcaacaacc	catgactgaa	ctggacagct	cctcgacct	3720
catctcttcc	ccagggaaga	agggggccgc	tcactctgac	cccagcaaga	cctctgtaga	3780
cacagggaaa	gtcagctggc	caggaatcc	cagccagcct	gcatcgccca	gggtcgccaa	3840
gtgcaaggcc	aggtctccag	tcaggctccc	ccatgagggc	agccccctcc	caggggagaa	3900
agcagcggct	ccccctgact	acagcaagac	tcgatcagca	tcggaaacca	gcacacccca	3960
caataccagg	agggtggctg	ccctcagggg	agcgggacct	ggagcagagg	gaatgacacc	4020
agctggtgct	gtcctgccag	gagacccct	cacatcccag	gagcagagac	agggagctcc	4080
aggtaaccac	agtaaggctc	tggaaatgac	aggaatccat	gcacctgaaa	gctcccagga	4140
gccttccctg	ctggagggag	cagattctgt	gtcctcaagg	gcaccgcagg	ccagcctctc	4200
catgctgcc	tccactgaca	acaccaaaga	agcatgtggc	catgtctcgg	ggcactgctg	4260
ccgggggggg	agtagagaga	gccctgtgac	ggacattgac	agcttcatca	aggagctgga	4320
tgcttctgca	gcaaggtctc	cgtcttccca	gacgggggac	agtggctctc	aggagggcag	4380
tgctcagggc	caccaccag	ccggggctgg	aggtgggagc	tcctgccgtg	ccgaaccagt	4440
ccgggggggg	cagacctcct	ccccgaggag	ggcctgggct	gctggtgcc	ccgcctaccc	4500
acaatgggcc	tcccagcctt	cggttttaga	ttcaattaat	cccgacaaac	attttactgt	4560
gaacaaaaac	tttctgagca	actactctag	aaatttttagc	agttttcatg	aagacagcac	4620
ctccctatca	ggcctgggtg	acagcacgga	gccgtctctg	tcatccatgt	atggcgatgc	4680
tgaggattct	tcttctgacc	ctgagtcact	cactgaagcc	ccacgagctt	ctgccaggga	4740
cggctggtcc	cctcctcggt	ccgctgtgtc	tttgcaaaag	gaagatcctt	cggagtcaga	4800
agaggaacag	attgagattt	gttccacacg	tggctgcccc	aatccaccct	cgagtctctg	4860
tcacttctcc	accagggctg	ccatctgtcc	tgctcagcc	aaagttctgt	cattaaaata	4920
cagcactccg	agagagtccg	tggccagtcc	ccgtgagaag	gtcgctgct	tgccaggctc	4980
atacattca	ggcccagact	cttcccagcc	atcatcactc	ttggagatga	gctctcagga	5040
gcatgaaact	catgcggaca	taagcacttc	acagaaccac	agggcctcgt	gtgcagaaga	5100
aaccacagaa	gtcaccagcg	ctagctcagc	catggaaaac	agtccgctgt	ctaaagtagc	5160
caggcatttt	cacagtccgc	ccatcattct	cagctcccc	aacatggtaa	atggcttgga	5220
acatgacctg	ctagatgacg	aaaccctgaa	tcaatacgaa	acaagcatta	atgcagctgc	5280
cagtctgtcc	tcttctagtg	tggatgtccc	taagaatgga	gaatctgttt	tggaaaacct	5340
ccacatctct	gaaagtcaag	acctggatga	cttgctacag	aaacaaaaa	tgatcgctag	5400
gaggcccatc	atggcctggt	ttaaagaaat	aaataaacat	aaccaaggca	cacatttgag	5460
gagcaaaacc	gagaaggaa	aaactcta	gcctgccaga	agtcccga	ccaagattca	5520
gatggtgagt	tcaagccaaa	aaaaggcg	tactgtgcct	catagccctc	ctcagccgaa	5580
aacaaacctg	gaaaataagg	acctgtctaa	gaagagtccg	gcagaaatgc	ttctgactaa	5640
tggtcagaag	gcaaagtgtg	gtccgaagct	gaagaggctc	agcctcaagg	gcaaggccaa	5700
agtcaactct	gaggccctg	ctgcgaatgc	tgtgaaggct	ggggggacgg	accacaggaa	5760
acccttgatc	tcacccca	cctcccacaa	aacactttct	aaggcagtgt	cacagcggct	5820
ccatgtagcc	gaccacgagg	acctgacag	aaacaccaca	gctgccccca	ggtcccccca	5880
gtgtgtgctg	gaaagcaagc	cacctcttgc	cacctctggg	ccactgaaac	cctcagtgtc	5940
tgacacgagc	atcaggacat	ttgtctcgcc	cctgacctct	cccaagcctg	ttcctgagca	6000
aggcatgtgg	agcaggttcc	acatggctgt	cctctctgaa	cccagacagag	gttgcccaac	6060

caccctaaa tctcctaagt gtagagcaga gggcagggcg ccccggtgctg actccggggc 6120
ggtgagtcgc gcagcgtcta ggaacggcat gtcccggtggca gggaacagac agagttagcc 6180
gcgcctggcc agccatgtgg cagcagacac agcccaaccc agggcgactg gcgaaaaagg 6240
aggcaacata atggccagcg atcgccctga aagaacaaac cagctgaaaa tctgtggagat 6300
ttctgctgaa gcagtgtcag agactgtatg tggttaacaag ccagctgaaa gcgacagacg 6360
gggaggggtgc ttggcccagg gcaactgtca ggagaagagt gaaatcaggc tctatcgcca 6420
ggtcgagaa tcacccacaa gtcacccatc ctcactccca tctcatgcct cccaggcaga 6480
gcaggaaatg tcacgatcat tcagcatggc aaaactggcg tctctctcct cctcccttca 6540
aacagccatt agaaaggcag aatactccca gggaaaatca agcctgatgt cagactcccg 6600
aggggtgccc agaaacagca ttccaggggg cccctcgggg gaggaccatc tctacttcac 6660
cccaaggcca gcgaccagga cctactccat gccagcccag ttctcaagcc attttggacg 6720
ggaggggtcac ccccccacac gcctgggtcg ctctcgggac agccaggctc ctgtgacaag 6780
cagtgttgtc cccgaggcaa aggcacccag aggtggtctt cccagcctgg ctaatggaca 6840
gggcatatat agtgtaaagc cgctgctgga cacatcgagg aatcttccag ccacagatga 6900
aggggatata atttcagtc aggagacgag ctgcctagtc acagacaaaa tcaaagtcac 6960
cagacgacac tactgctatg agcagaactg gccccatgaa tctacctcat ttttctctgt 7020
gaagcagcgg atcaagtctt ttgagaacct ggccaatgct gaccggcctg tagccaagtc 7080
cggggcttcc ccatttttgt cgggtgagctc caagcctccc attgggaggc ggtcttccgg 7140
cagcattgtt tccgggagcc tgggcccacc aggtgacgca gcagcaaggc tggtgagacg 7200
cagcttgagt tctgcagcg aaaaccaaag cgaagccggc accctcctgc cccagatggc 7260
caagtctccc tcaatcatga cactgaccat ctctcggcag aaccaccag agaccagtag 7320
caagggtctt gattcggaac taaagaaatc acttggtcct ttgggaattc ccacccaac 7380
gatgaccctg gcttctcctg ttaagaggaa caagtcctcg gtacgccaca cgcagccctc 7440
gcccggtgct cgctccaagc tccaggagct gagagccttg agcatgcctg accttgacaa 7500
gctctgcagc gaggattact cagcagggcc gagcgccgtg ctcttcaaaa ctgagctgga 7560
gatcaccccc aggaggtcac ctggccctcc tgctggaggc gtttctgtgc ccgagaaggg 7620
cgggaacagg gcctgtccag gaggaagtgg ccctaaacc agtgctgctg agacaccag 7680
ttcagccagt gatacgggtg aagctgcccc ggatctgcct tttagaagaa gctggtcagt 7740
taatttggat caacttctag tctcagcggg ggaccagcaa agattacagt ctgttttatc 7800
gtcagtggga tcgaaatcta ccacccaaac tctcattcag gaagcgaaag cacaatcaga 7860
gaatgaagaa gatgtttgct tcatagtctt gaatagaaaa gaaggctcag gtctgggatt 7920
cagtgtggca ggagggacag atgtggagcc aaaatcaatc acggtccaca ggggtgtttc 7980
tcagggggcg gcttctcagg aagggactat gaaccgaggg gatttccttc tgtcagtcaa 8040
cggcgcccca ctggctggct tagcccacgg gaatgtcctg aaggttctgc accaggcaca 8100
gctgcacaaa gatgcctcg tggatcatca gaaagggatg gatcagccca ggccctctgc 8160
ccggcaggag cctcccacag ccaatgggaa gggtttgctg tccagaaaga ccatccccc 8220
ggagcctggc attgggagaa gtgtggctgt acacgatgct ctgtgtgttg aagtgtgaa 8280
gacctcggct gggctgggac tgaatctgga tgggggaaaa tcatcgggtg cgggagatgg 8340
gcccttggtc attaaaagag tgtacaaagg tgggtcggct gaacaagctg gaataataga 8400
agctggagat gaaattcttg ctattaatgg gaaacctctg gttgggctca tgcactttga 8460
tgcttgaat attatgaagt ctgtcccaga aggacctgtg cagttattaa ttagaaagca 8520
taggaattct tcatgaattt taacaagaat cattttctca gttctcttct ttcttttagca 8580
aatcagagtg acttctttta accacagggt gttgaaatgg ccaacactgg tacagacagc 8640

<210> 27

<211> 2811

<212> PRT

<213> Homo sapiens

<400> 27

Met	Pro	Ile	Thr	Gln	Asp	Asn	Ala	Val	Leu	His	Leu	Pro	Leu	Leu	Tyr
1				5					10					15	

Gln	Trp	Leu	Gln	Asn	Ser	Leu	Gln	Glu	Gly	Gly	Asp	Gly	Pro	Glu	Gln
			20					25					30		

Arg Leu Cys Gln Ala Ala Ile Gln Lys Leu Gln Glu Tyr Ile Gln Leu
 35 40 45
 Asn Phe Ala Val Asp Glu Ser Thr Val Pro Pro Asp His Ser Pro Pro
 50 55 60
 Glu Met Glu Ile Cys Thr Val Tyr Leu Thr Lys Glu Leu Gly Asp Thr
 65 70 75 80
 Glu Thr Val Gly Leu Ser Phe Gly Asn Ile Pro Val Phe Gly Asp Tyr
 85 90 95
 Gly Glu Lys Arg Arg Gly Gly Lys Lys Arg Lys Thr His Gln Gly Pro
 100 105 110
 Val Leu Asp Val Gly Cys Ile Trp Val Thr Glu Leu Arg Lys Asn Ser
 115 120 125
 Pro Ala Gly Lys Ser Gly Lys Val Arg Leu Arg Asp Glu Ile Leu Ser
 130 135 140
 Leu Asn Gly Gln Leu Met Val Gly Val Asp Val Ser Gly Ala Ser Tyr
 145 150 155 160
 Leu Ala Glu Gln Cys Trp Asn Gly Gly Phe Ile Tyr Leu Ile Met Leu
 165 170 175
 Arg Arg Phe Lys His Lys Ala His Ser Thr Tyr Asn Gly Asn Ser Ser
 180 185 190
 Asn Ser Ser Glu Pro Gly Glu Thr Pro Thr Leu Glu Leu Gly Asp Arg
 195 200 205
 Thr Ala Lys Lys Gly Lys Arg Thr Arg Lys Phe Gly Val Ile Ser Arg
 210 215 220
 Pro Pro Ala Asn Lys Ala Pro Glu Glu Ser Lys Gly Ser Ala Gly Cys
 225 230 235 240
 Glu Val Ser Ser Asp Pro Ser Thr Glu Leu Glu Asn Gly Leu Asp Pro
 245 250 255
 Glu Leu Gly Asn Gly His Val Phe Gln Leu Glu Asn Gly Pro Asp Ser
 260 265 270
 Leu Lys Glu Val Ala Gly Pro His Leu Glu Arg Ser Glu Val Asp Arg
 275 280 285
 Gly Thr Glu His Arg Ile Pro Lys Thr Asp Ala Pro Leu Thr Thr Ser
 290 295 300
 Asn Asp Lys Arg Arg Phe Ser Lys Gly Gly Lys Thr Asp Phe Gln Ser
 305 310 315 320
 Ser Asp Cys Leu Ala Arg Ser Lys Glu Glu Val Gly Arg Ile Trp Lys
 325 330 335

Met Glu Leu Leu Lys Glu Ser Asp Gly Leu Gly Ile Gln Val Ser Gly
 340 345 350
 Gly Arg Gly Ser Lys Arg Ser Pro His Ala Ile Val Val Thr Gln Val
 355 360 365
 Lys Glu Gly Gly Ala Ala His Arg Leu Arg Asp Gly Arg Leu Ser Leu
 370 375 380
 Gly Asp Glu Leu Leu Val Ile Asn Gly His Leu Leu Val Gly Leu Ser
 385 390 395 400
 His Glu Glu Ala Val Ala Ile Leu Arg Ser Ala Thr Gly Met Val Gln
 405 410 415
 Leu Val Val Ala Ser Lys Val Gly Val Leu Ser Ala Phe Gln Met Pro
 420 425 430
 Gly Thr Asp Glu Pro Gln Asp Val Cys Gly Ala Glu Glu Ser Lys Gly
 435 440 445
 Asn Leu Glu Ser Pro Lys Gln Gly Ser Asn Lys Ile Lys Leu Lys Ser
 450 455 460
 Arg Leu Ser Gly Arg Trp Gly Leu Tyr Leu Met Gln Pro Val Gly Gly
 465 470 475 480
 Val His Arg Leu Glu Ser Val Glu Glu Tyr Asn Glu Leu Met Val Arg
 485 490 495
 Asn Gly Asp Pro Arg Ile Arg Met Leu Glu Val Ser Arg Asp Gly Arg
 500 505 510
 Lys His Ser Leu Pro Gln Leu Leu Asp Ser Ser Ser Ala Ser Gln Glu
 515 520 525
 Tyr His Ile Val Lys Lys Ser Thr Arg Ser Leu Ser Thr Thr Gln Val
 530 535 540
 Glu Ser Pro Arg Arg Leu Ile Arg Pro Ser Val Ile Ser Ile Ile Gly
 545 550 555 560
 Leu Tyr Lys Glu Lys Gly Lys Gly Leu Gly Phe Ser Ile Ala Gly Gly
 565 570 575
 Arg Asp Cys Ile Arg Gly Gln Met Gly Ile Phe Val Lys Thr Ile Phe
 580 585 590
 Pro Asn Gly Ser Ala Ala Glu Asp Gly Arg Leu Lys Glu Gly Asp Glu
 595 600 605
 Ile Leu Asp Val Asn Gly Ile Pro Ile Lys Gly Leu Thr Phe Gln Glu
 610 615 620
 Ala Ile His Thr Phe Lys Gln Ile Arg Ser Gly Leu Phe Val Leu Thr
 625 630 635 640

Val Arg Thr Lys Leu Val Ser Pro Ser Leu Thr Pro Cys Ser Thr Pro
645 650 655
Thr His Met Ser Arg Ser Ala Ser Pro Asn Phe Asn Thr Ser Gly Gly
660 665 670
Ala Ser Ala Gly Gly Ser Asp Glu Gly Ser Ser Ser Ser Leu Gly Arg
675 680 685
Lys Thr Pro Gly Pro Lys Asp Arg Ile Val Met Glu Val Thr Leu Asn
690 695 700
Lys Glu Pro Arg Val Gly Leu Gly Ile Gly Ala Cys Cys Leu Ala Leu
705 710 715 720
Glu Asn Ser Pro Pro Gly Ile Tyr Ile His Ser Leu Ala Pro Gly Ser
725 730 735
Val Ala Lys Met Glu Ser Asn Leu Ser Arg Gly Asp Gln Ile Leu Glu
740 745 750
Val Asn Ser Val Asn Val Arg His Ala Ala Leu Ser Lys Val His Ala
755 760 765
Ile Leu Ser Lys Cys Pro Pro Gly Pro Val Arg Leu Val Ile Gly Arg
770 775 780
His Pro Asn Pro Lys Val Ser Glu Gln Glu Met Asp Glu Val Ile Ala
785 790 795 800
Arg Ser Thr Tyr Gln Glu Ser Lys Glu Ala Asn Ser Ser Pro Gly Leu
805 810 815
Gly Thr Pro Leu Lys Ser Pro Ser Leu Ala Lys Lys Asp Ser Leu Ile
820 825 830
Ser Glu Ser Glu Leu Ser Gln Tyr Phe Ala His Asp Val Pro Gly Pro
835 840 845
Leu Ser Asp Phe Met Val Val Gly Ser Glu Asp Glu Asp His Pro Gly
850 855 860
Ser Gly Cys Ser Thr Ser Glu Glu Gly Ser Leu Pro Pro Ser Thr Ser
865 870 875 880
Thr His Lys Glu Pro Gly Lys Pro Arg Ala Asn Ser Leu Val Thr Leu
885 890 895
Gly Ser His Arg Ala Ser Gly Leu Phe His Lys Gln Val Thr Val Ala
900 905 910
Arg Gln Ala Ser Leu Pro Gly Ser Pro Gln Ala Leu Arg Asn Pro Leu
915 920 925
Leu Arg Gln Arg Lys Val Gly Cys Tyr Asp Ala Asn Asp Ala Ser Asp
930 935 940

Glu Glu Glu Phe Asp Arg Glu Gly Asp Cys Ile Ser Leu Pro Gly Ala
 945 950 955 960
 Leu Pro Gly Pro Ile Arg Pro Leu Ser Glu Asp Asp Pro Arg Arg Val
 965 970 975
 Ser Ile Ser Ser Ser Lys Gly Met Asp Val His Asn Gln Glu Glu Arg
 980 985 990
 Pro Arg Lys Thr Leu Val Ser Lys Ala Ile Ser Ala Pro Leu Leu Gly
 995 1000 1005
 Ser Ser Val Asp Leu Glu Glu Ser Ile Pro Glu Gly Met Val Asp Ala
 1010 1015 1020
 Ala Ser Tyr Ala Ala Asn Leu Thr Asp Ser Ala Glu Ala Pro Lys Gly
 1025 1030 1035 1040
 Ser Pro Gly Ser Trp Trp Lys Lys Glu Leu Ser Gly Ser Ser Ser Ala
 1045 1050 1055
 Pro Lys Leu Glu Tyr Thr Val Arg Thr Asp Thr Gln Ser Pro Thr Asn
 1060 1065 1070
 Thr Gly Ser Pro Ser Ser Pro Gln Gln Lys Ser Glu Gly Leu Gly Ser
 1075 1080 1085
 Arg His Arg Pro Val Ala Arg Val Ser Pro His Cys Lys Arg Ser Glu
 1090 1095 1100
 Ala Glu Ala Lys Pro Ser Gly Ser Gln Thr Val Asn Leu Thr Gly Arg
 1105 1110 1115 1120
 Ala Asn Asp Pro Cys Asp Leu Asp Ser Arg Val Gln Ala Thr Ser Val
 1125 1130 1135
 Lys Val Thr Val Ala Gly Phe Gln Pro Gly Gly Ala Val Glu Lys Glu
 1140 1145 1150
 Ser Leu Gly Lys Leu Thr Thr Gly Asp Ala Cys Val Ser Thr Ser Cys
 1155 1160 1165
 Glu Leu Ala Ser Ala Leu Ser His Leu Asp Ala Ser His Leu Thr Glu
 1170 1175 1180
 Asn Leu Pro Lys Ala Ala Ser Glu Leu Gly Gln Gln Pro Met Thr Glu
 1185 1190 1195 1200
 Leu Asp Ser Ser Ser Asp Leu Ile Ser Ser Pro Gly Lys Lys Gly Ala
 1205 1210 1215
 Ala His Pro Asp Pro Ser Lys Thr Ser Val Asp Thr Gly Lys Val Ser
 1220 1225 1230
 Arg Pro Glu Asn Pro Ser Gln Pro Ala Ser Pro Arg Val Ala Lys Cys
 1235 1240 1245

Lys Ala Arg Ser Pro Val Arg Leu Pro His Glu Gly Ser Pro Ser Pro
 1250 1255 1260

Gly Glu Lys Ala Ala Ala Pro Pro Asp Tyr Ser Lys Thr Arg Ser Ala
 1265 1270 1275 1280

Ser Glu Thr Ser Thr Pro His Asn Thr Arg Arg Val Ala Ala Leu Arg
 1285 1290 1295

Gly Ala Gly Pro Gly Ala Glu Gly Met Thr Pro Ala Gly Ala Val Leu
 1300 1305 1310

Pro Gly Asp Pro Leu Thr Ser Gln Glu Gln Arg Gln Gly Ala Pro Gly
 1315 1320 1325

Asn His Ser Lys Ala Leu Glu Met Thr Gly Ile His Ala Pro Glu Ser
 1330 1335 1340

Ser Gln Glu Pro Ser Leu Leu Glu Gly Ala Asp Ser Val Ser Ser Arg
 1345 1350 1355 1360

Ala Pro Gln Ala Ser Leu Ser Met Leu Pro Ser Thr Asp Asn Thr Lys
 1365 1370 1375

Glu Ala Cys Gly His Val Ser Gly His Cys Cys Pro Gly Gly Ser Arg
 1380 1385 1390

Glu Ser Pro Val Thr Asp Ile Asp Ser Phe Ile Lys Glu Leu Asp Ala
 1395 1400 1405

Ser Ala Ala Arg Ser Pro Ser Ser Gln Thr Gly Asp Ser Gly Ser Gln
 1410 1415 1420

Glu Gly Ser Ala Gln Gly His Pro Pro Ala Gly Ala Gly Gly Gly Ser
 1425 1430 1435 1440

Ser Cys Arg Ala Glu Pro Val Pro Gly Gly Gln Thr Ser Ser Pro Arg
 1445 1450 1455

Arg Ala Trp Ala Ala Gly Ala Pro Ala Tyr Pro Gln Trp Ala Ser Gln
 1460 1465 1470

Pro Ser Val Leu Asp Ser Ile Asn Pro Asp Lys His Phe Thr Val Asn
 1475 1480 1485

Lys Asn Phe Leu Ser Asn Tyr Ser Arg Asn Phe Ser Ser Phe His Glu
 1490 1495 1500

Asp Ser Thr Ser Leu Ser Gly Leu Gly Asp Ser Thr Glu Pro Ser Leu
 1505 1510 1515 1520

Ser Ser Met Tyr Gly Asp Ala Glu Asp Ser Ser Ser Asp Pro Glu Ser
 1525 1530 1535

Leu Thr Glu Ala Pro Arg Ala Ser Ala Arg Asp Gly Trp Ser Pro Pro
 1540 1545 1550

Arg Ser Arg Val Ser Leu His Lys Glu Asp Pro Ser Glu Ser Glu Glu
 1555 1560 1565
 Glu Gln Ile Glu Ile Cys Ser Thr Arg Gly Cys Pro Asn Pro Pro Ser
 1570 1575 1580
 Ser Pro Ala His Leu Pro Thr Gln Ala Ala Ile Cys Pro Ala Ser Ala
 1585 1590 1595 1600
 Lys Val Leu Ser Leu Lys Tyr Ser Thr Pro Arg Glu Ser Val Ala Ser
 1605 1610 1615
 Pro Arg Glu Lys Val Ala Cys Leu Pro Gly Ser Tyr Thr Ser Gly Pro
 1620 1625 1630
 Asp Ser Ser Gln Pro Ser Ser Leu Leu Glu Met Ser Ser Gln Glu His
 1635 1640 1645
 Glu Thr His Ala Asp Ile Ser Thr Ser Gln Asn His Arg Pro Ser Cys
 1650 1655 1660
 Ala Glu Glu Thr Thr Glu Val Thr Ser Ala Ser Ser Ala Met Glu Asn
 1665 1670 1675 1680
 Ser Pro Leu Ser Lys Val Ala Arg His Phe His Ser Pro Pro Ile Ile
 1685 1690 1695
 Leu Ser Ser Pro Asn Met Val Asn Gly Leu Glu His Asp Leu Leu Asp
 1700 1705 1710
 Asp Glu Thr Leu Asn Gln Tyr Glu Thr Ser Ile Asn Ala Ala Ala Ser
 1715 1720 1725
 Leu Ser Ser Phe Ser Val Asp Val Pro Lys Asn Gly Glu Ser Val Leu
 1730 1735 1740
 Glu Asn Leu His Ile Ser Glu Ser Gln Asp Leu Asp Asp Leu Leu Gln
 1745 1750 1755 1760
 Lys Pro Lys Met Ile Ala Arg Arg Pro Ile Met Ala Trp Phe Lys Glu
 1765 1770 1775
 Ile Asn Lys His Asn Gln Gly Thr His Leu Arg Ser Lys Thr Glu Lys
 1780 1785 1790
 Glu Gln Pro Leu Met Pro Ala Arg Ser Pro Asp Ser Lys Ile Gln Met
 1795 1800 1805
 Val Ser Ser Ser Gln Lys Lys Gly Val Thr Val Pro His Ser Pro Pro
 1810 1815 1820
 Gln Pro Lys Thr Asn Leu Glu Asn Lys Asp Leu Ser Lys Lys Ser Pro
 1825 1830 1835 1840
 Ala Glu Met Leu Leu Thr Asn Gly Gln Lys Ala Lys Cys Gly Pro Lys
 1845 1850 1855

Leu Lys Arg Leu Ser Leu Lys Gly Lys Ala Lys Val Asn Ser Glu Ala
 1860 1865 1870
 Pro Ala Ala Asn Ala Val Lys Ala Gly Gly Thr Asp His Arg Lys Pro
 1875 1880 1885
 Leu Ile Ser Pro Gln Thr Ser His Lys Thr Leu Ser Lys Ala Val Ser
 1890 1895 1900
 Gln Arg Leu His Val Ala Asp His Glu Asp Pro Asp Arg Asn Thr Thr
 1905 1910 1915 1920
 Ala Ala Pro Arg Ser Pro Gln Cys Val Leu Glu Ser Lys Pro Pro Leu
 1925 1930 1935
 Ala Thr Ser Gly Pro Leu Lys Pro Ser Val Ser Asp Thr Ser Ile Arg
 1940 1945 1950
 Thr Phe Val Ser Pro Leu Thr Ser Pro Lys Pro Val Pro Glu Gln Gly
 1955 1960 1965
 Met Trp Ser Arg Phe His Met Ala Val Leu Ser Glu Pro Asp Arg Gly
 1970 1975 1980
 Cys Pro Thr Thr Pro Lys Ser Pro Lys Cys Arg Ala Glu Gly Arg Ala
 1985 1990 1995 2000
 Pro Arg Ala Asp Ser Gly Pro Val Ser Pro Ala Ala Ser Arg Asn Gly
 2005 2010 2015
 Met Ser Val Ala Gly Asn Arg Gln Ser Glu Pro Arg Leu Ala Ser His
 2020 2025 2030
 Val Ala Ala Asp Thr Ala Gln Pro Arg Pro Thr Gly Glu Lys Gly Gly
 2035 2040 2045
 Asn Ile Met Ala Ser Asp Arg Leu Glu Arg Thr Asn Gln Leu Lys Ile
 2050 2055 2060
 Val Glu Ile Ser Ala Glu Ala Val Ser Glu Thr Val Cys Gly Asn Lys
 2065 2070 2075 2080
 Pro Ala Glu Ser Asp Arg Arg Gly Gly Cys Leu Ala Gln Gly Asn Cys
 2085 2090 2095
 Gln Glu Lys Ser Glu Ile Arg Leu Tyr Arg Gln Val Ala Glu Ser Ser
 2100 2105 2110
 Thr Ser His Pro Ser Ser Leu Pro Ser His Ala Ser Gln Ala Glu Gln
 2115 2120 2125
 Glu Met Ser Arg Ser Phe Ser Met Ala Lys Leu Ala Ser Ser Ser Ser
 2130 2135 2140
 Ser Leu Gln Thr Ala Ile Arg Lys Ala Glu Tyr Ser Gln Gly Lys Ser
 2145 2150 2155 2160

Ser Leu Met Ser Asp Ser Arg Gly Val Pro Arg Asn Ser Ile Pro Gly
 2165 2170 2175
 Gly Pro Ser Gly Glu Asp His Leu Tyr Phe Thr Pro Arg Pro Ala Thr
 2180 2185 2190
 Arg Thr Tyr Ser Met Pro Ala Gln Phe Ser Ser His Phe Gly Arg Glu
 2195 2200 2205
 Gly His Pro Pro His Ser Leu Gly Arg Ser Arg Asp Ser Gln Val Pro
 2210 2215 2220
 Val Thr Ser Ser Val Val Pro Glu Ala Lys Ala Ser Arg Gly Gly Leu
 2225 2230 2235 2240
 Pro Ser Leu Ala Asn Gly Gln Gly Ile Tyr Ser Val Lys Pro Leu Leu
 2245 2250 2255
 Asp Thr Ser Arg Asn Leu Pro Ala Thr Asp Glu Gly Asp Ile Ile Ser
 2260 2265 2270
 Val Gln Glu Thr Ser Cys Leu Val Thr Asp Lys Ile Lys Val Thr Arg
 2275 2280 2285
 Arg His Tyr Cys Tyr Glu Gln Asn Trp Pro His Glu Ser Thr Ser Phe
 2290 2295 2300
 Phe Ser Val Lys Gln Arg Ile Lys Ser Phe Glu Asn Leu Ala Asn Ala
 2305 2310 2315 2320
 Asp Arg Pro Val Ala Lys Ser Gly Ala Ser Pro Phe Leu Ser Val Ser
 2325 2330 2335
 Ser Lys Pro Pro Ile Gly Arg Arg Ser Ser Gly Ser Ile Val Ser Gly
 2340 2345 2350
 Ser Leu Gly His Pro Gly Asp Ala Ala Ala Arg Leu Leu Arg Arg Ser
 2355 2360 2365
 Leu Ser Ser Cys Ser Glu Asn Gln Ser Glu Ala Gly Thr Leu Leu Pro
 2370 2375 2380
 Gln Met Ala Lys Ser Pro Ser Ile Met Thr Leu Thr Ile Ser Arg Gln
 2385 2390 2395 2400
 Asn Pro Pro Glu Thr Ser Ser Lys Gly Ser Asp Ser Glu Leu Lys Lys
 2405 2410 2415
 Ser Leu Gly Pro Leu Gly Ile Pro Thr Pro Thr Met Thr Leu Ala Ser
 2420 2425 2430
 Pro Val Lys Arg Asn Lys Ser Ser Val Arg His Thr Gln Pro Ser Pro
 2435 2440 2445
 Val Ser Arg Ser Lys Leu Gln Glu Leu Arg Ala Leu Ser Met Pro Asp
 2450 2455 2460

Leu Asp Lys Leu Cys Ser Glu Asp Tyr Ser Ala Gly Pro Ser Ala Val
 2465 2470 2475 2480

Leu Phe Lys Thr Glu Leu Glu Ile Thr Pro Arg Arg Ser Pro Gly Pro
 2485 2490 2495

Pro Ala Gly Gly Val Ser Cys Pro Glu Lys Gly Gly Asn Arg Ala Cys
 2500 2505 2510

Pro Gly Gly Ser Gly Pro Lys Thr Ser Ala Ala Glu Thr Pro Ser Ser
 2515 2520 2525

Ala Ser Asp Thr Gly Glu Ala Ala Gln Asp Leu Pro Phe Arg Arg Ser
 2530 2535 2540

Trp Ser Val Asn Leu Asp Gln Leu Leu Val Ser Ala Gly Asp Gln Gln
 2545 2550 2555 2560

Arg Leu Gln Ser Val Leu Ser Ser Val Gly Ser Lys Ser Thr Ile Leu
 2565 2570 2575

Thr Leu Ile Gln Glu Ala Lys Ala Gln Ser Glu Asn Glu Glu Asp Val
 2580 2585 2590

Cys Phe Ile Val Leu Asn Arg Lys Glu Gly Ser Gly Leu Gly Phe Ser
 2595 2600 2605

Val Ala Gly Gly Thr Asp Val Glu Pro Lys Ser Ile Thr Val His Arg
 2610 2615 2620

Val Phe Ser Gln Gly Ala Ala Ser Gln Glu Gly Thr Met Asn Arg Gly
 2625 2630 2635 2640

Asp Phe Leu Leu Ser Val Asn Gly Ala Ser Leu Ala Gly Leu Ala His
 2645 2650 2655

Gly Asn Val Leu Lys Val Leu His Gln Ala Gln Leu His Lys Asp Ala
 2660 2665 2670

Leu Val Val Ile Lys Lys Gly Met Asp Gln Pro Arg Pro Ser Ala Arg
 2675 2680 2685

Gln Glu Pro Pro Thr Ala Asn Gly Lys Gly Leu Leu Ser Arg Lys Thr
 2690 2695 2700

Ile Pro Leu Glu Pro Gly Ile Gly Arg Ser Val Ala Val His Asp Ala
 2705 2710 2715 2720

Leu Cys Val Glu Val Leu Lys Thr Ser Ala Gly Leu Gly Leu Ser Leu
 2725 2730 2735

Asp Gly Gly Lys Ser Ser Val Thr Gly Asp Gly Pro Leu Val Ile Lys
 2740 2745 2750

Arg Val Tyr Lys Gly Gly Ala Ala Glu Gln Ala Gly Ile Ile Glu Ala
 2755 2760 2765

Gly Asp Glu Ile Leu Ala Ile Asn Gly Lys Pro Leu Val Gly Leu Met
 2770 2775 2780

His Phe Asp Ala Trp Asn Ile Met Lys Ser Val Pro Glu Gly Pro Val
 2785 2790 2795 2800

Gln Leu Leu Ile Arg Lys His Arg Asn Ser Ser
 2805 2810

<210> 28

<211> 798

<212> PRT

<213> Homo sapiens

<400> 28

Met Leu Ile Asn Cys Glu Ala Lys Gly Ile Lys Met Val Ser Glu Ile
 1 5 10 15

Ser Val Pro Pro Ser Arg Pro Phe Gln Leu Ser Leu Leu Asn Asn Gly
 20 25 30

Leu Thr Met Leu His Thr Asn Asp Phe Ser Gly Leu Thr Asn Ala Ile
 35 40 45

Ser Ile His Leu Gly Phe Asn Asn Ile Ala Asp Ile Glu Ile Gly Ala
 50 55 60

Phe Asn Gly Leu Gly Leu Leu Lys Gln Leu His Ile Asn His Asn Ser
 65 70 75 80

Leu Glu Ile Leu Lys Glu Asp Thr Phe His Gly Leu Glu Asn Leu Glu
 85 90 95

Phe Leu Gln Ala Asp Asn Asn Phe Ile Thr Val Ile Glu Pro Ser Ala
 100 105 110

Phe Ser Lys Leu Asn Arg Leu Lys Val Leu Ile Leu Asn Asp Asn Ala
 115 120 125

Ile Glu Ser Leu Pro Pro Asn Ile Phe Arg Phe Val Pro Leu Thr His
 130 135 140

Leu Asp Leu Arg Gly Asn Gln Leu Gln Thr Leu Pro Tyr Val Gly Phe
 145 150 155 160

Leu Glu His Ile Gly Arg Ile Leu Asp Leu Gln Leu Glu Asp Asn Lys
 165 170 175

Trp Ala Cys Asn Cys Asp Leu Leu Gln Leu Lys Thr Trp Leu Glu Asn
 180 185 190

Met Pro Pro Gln Ser Ile Ile Gly Asp Val Val Cys Asn Ser Pro Pro
 195 200 205

Phe Phe Lys Gly Ser Ile Leu Ser Arg Leu Lys Lys Glu Ser Ile Cys
 210 215 220

Pro Thr Pro Pro Val Tyr Glu Glu His Glu Asp Pro Ser Gly Ser Leu
 225 230 235 240
 His Leu Ala Ala Thr Ser Ser Ile Asn Asp Ser Arg Met Ser Thr Lys
 245 250 255
 Thr Thr Ser Ile Leu Lys Leu Pro Thr Lys Ala Pro Gly Leu Ile Pro
 260 265 270
 Tyr Ile Thr Lys Pro Ser Thr Gln Leu Pro Gly Pro Tyr Cys Pro Ile
 275 280 285
 Pro Cys Asn Cys Lys Val Leu Ser Pro Ser Gly Leu Leu Ile His Cys
 290 295 300
 Gln Glu Arg Asn Ile Glu Ser Leu Ser Asp Leu Arg Pro Pro Pro Gln
 305 310 315 320
 Asn Pro Arg Lys Leu Ile Leu Ala Gly Asn Ile Ile His Ser Leu Met
 325 330 335
 Lys Ser Asp Leu Val Glu Tyr Phe Thr Leu Glu Met Leu His Leu Gly
 340 345 350
 Asn Asn Arg Ile Glu Val Leu Glu Glu Gly Ser Phe Met Asn Leu Thr
 355 360 365
 Arg Leu Gln Lys Leu Tyr Leu Asn Gly Asn His Leu Thr Lys Leu Ser
 370 375 380
 Lys Gly Met Phe Leu Gly Leu His Asn Leu Glu Tyr Leu Tyr Leu Glu
 385 390 395 400
 Tyr Asn Ala Ile Lys Glu Ile Leu Pro Gly Thr Phe Asn Pro Met Pro
 405 410 415
 Lys Leu Lys Val Leu Tyr Leu Asn Asn Asn Leu Leu Gln Val Leu Pro
 420 425 430
 Pro His Ile Phe Ser Gly Val Pro Leu Thr Lys Val Asn Leu Lys Thr
 435 440 445
 Asn Gln Phe Thr His Leu Pro Val Ser Asn Ile Leu Asp Asp Leu Asp
 450 455 460
 Leu Leu Thr Gln Ile Asp Leu Glu Asp Asn Pro Trp Asp Cys Ser Cys
 465 470 475 480
 Asp Leu Val Gly Leu Gln Gln Trp Ile Gln Lys Leu Ser Lys Asn Thr
 485 490 495
 Val Thr Asp Asp Ile Leu Cys Thr Ser Pro Gly His Leu Asp Lys Lys
 500 505 510
 Glu Leu Lys Ala Leu Asn Ser Glu Ile Leu Cys Pro Gly Leu Val Asn
 515 520 525

Asn Pro Ser Met Pro Thr Gln Thr Ser Tyr Leu Met Val Thr Thr Pro
 530 535 540
 Ala Thr Thr Thr Asn Thr Ala Asp Thr Ile Leu Arg Ser Leu Thr Asp
 545 550 555 560
 Ala Val Pro Leu Ser Val Leu Ile Leu Gly Leu Leu Ile Met Phe Ile
 565 570 575
 Thr Ile Val Phe Cys Ala Ala Gly Ile Val Val Leu Val Leu His Arg
 580 585 590
 Arg Arg Arg Tyr Lys Lys Lys Gln Val Asp Glu Gln Met Arg Asp Asn
 595 600 605
 Ser Pro Val His Leu Gln Tyr Ser Met Tyr Gly His Lys Thr Thr His
 610 615 620
 His Thr Thr Glu Arg Pro Ser Ala Ser Leu Tyr Glu Gln His Met Val
 625 630 635 640
 Ser Pro Met Val His Val Tyr Arg Ser Pro Ser Phe Gly Pro Lys His
 645 650 655
 Leu Glu Glu Glu Glu Glu Arg Asn Glu Lys Glu Gly Ser Asp Ala Lys
 660 665 670
 His Leu Gln Arg Ser Leu Leu Glu Gln Glu Asn His Ser Pro Leu Thr
 675 680 685
 Gly Ser Asn Met Lys Tyr Lys Thr Thr Asn Gln Ser Thr Glu Phe Leu
 690 695 700
 Ser Phe Gln Asp Ala Ser Ser Leu Tyr Arg Asn Ile Leu Glu Lys Glu
 705 710 715 720
 Arg Glu Leu Gln Gln Leu Gly Ile Thr Glu Tyr Leu Arg Lys Asn Ile
 725 730 735
 Ala Gln Leu Gln Pro Asp Met Glu Ala His Tyr Pro Gly Ala His Glu
 740 745 750
 Glu Leu Lys Leu Met Glu Thr Leu Met Tyr Ser Arg Pro Arg Lys Val
 755 760 765
 Leu Val Glu Gln Thr Lys Asn Glu Tyr Phe Glu Leu Lys Ala Asn Leu
 770 775 780
 His Ala Glu Pro Asp Tyr Leu Glu Val Leu Glu Gln Gln Thr
 785 790 795

<210> 29
 <211> 440
 <212> PRT
 <213> Homo sapiens

<400> 29

Met	Leu	Ile	Asn	Cys	Glu	Ala	Lys	Gly	Ile	Lys	Met	Val	Ser	Glu	Ile	1	5	10	15
Ser	Val	Pro	Pro	Ser	Arg	Pro	Phe	Gln	Leu	Ser	Leu	Leu	Asn	Asn	Gly	20	25	30	
Leu	Thr	Met	Leu	His	Thr	Asn	Asp	Phe	Ser	Gly	Leu	Thr	Asn	Ala	Ile	35	40	45	
Ser	Ile	His	Leu	Gly	Phe	Asn	Asn	Ile	Ala	Asp	Ile	Glu	Ile	Gly	Ala	50	55	60	
Phe	Asn	Gly	Leu	Gly	Leu	Leu	Lys	Gln	Leu	His	Ile	Asn	His	Asn	Ser	65	70	75	80
Leu	Glu	Ile	Leu	Lys	Glu	Asp	Thr	Phe	His	Gly	Leu	Glu	Asn	Leu	Glu	85	90	95	
Phe	Leu	Gln	Ala	Asp	Asn	Asn	Phe	Ile	Thr	Val	Ile	Glu	Pro	Ser	Ala	100	105	110	
Phe	Ser	Lys	Leu	Asn	Arg	Leu	Lys	Val	Leu	Ile	Leu	Asn	Asp	Asn	Ala	115	120	125	
Ile	Glu	Ser	Leu	Pro	Pro	Asn	Ile	Phe	Arg	Phe	Val	Pro	Leu	Thr	His	130	135	140	
Leu	Asp	Leu	Arg	Gly	Asn	Gln	Leu	Gln	Thr	Leu	Pro	Tyr	Val	Gly	Phe	145	150	155	160
Leu	Glu	His	Ile	Gly	Arg	Ile	Leu	Asp	Leu	Gln	Leu	Glu	Asp	Asn	Lys	165	170	175	
Trp	Ala	Cys	Asn	Cys	Asp	Leu	Leu	Gln	Leu	Lys	Thr	Trp	Leu	Glu	Asn	180	185	190	
Met	Pro	Pro	Gln	Ser	Ile	Ile	Gly	Asp	Val	Val	Cys	Asn	Ser	Pro	Pro	195	200	205	
Phe	Phe	Lys	Gly	Ser	Ile	Leu	Ser	Arg	Leu	Lys	Lys	Glu	Ser	Ile	Cys	210	215	220	
Pro	Thr	Pro	Pro	Val	Tyr	Glu	Glu	His	Glu	Asp	Pro	Ser	Gly	Ser	Leu	225	230	235	240
His	Leu	Ala	Ala	Thr	Ser	Ser	Ile	Asn	Asp	Ser	Arg	Met	Ser	Thr	Lys	245	250	255	
Thr	Thr	Ser	Ile	Leu	Lys	Leu	Pro	Thr	Lys	Ala	Pro	Gly	Leu	Ile	Pro	260	265	270	
Tyr	Ile	Thr	Lys	Pro	Ser	Thr	Gln	Leu	Pro	Gly	Pro	Tyr	Cys	Pro	Ile	275	280	285	
Pro	Cys	Asn	Cys	Lys	Val	Leu	Ser	Pro	Ser	Gly	Leu	Leu	Ile	His	Cys				

Asn Ile Glu Gly Gly Ala Phe Leu Gly Leu Ser Ala Leu Lys Gln Leu
 115 120 125
 His Leu Asn Asn Asn Glu Leu Lys Ile Leu Arg Ala Asp Thr Phe Leu
 130 135 140
 Gly Ile Glu Asn Leu Glu Tyr Leu Gln Ala Asp Tyr Asn Leu Ile Lys
 145 150 155 160
 Tyr Ile Glu Arg Gly Ala Phe Asn Lys Leu His Lys Leu Lys Val Leu
 165 170 175
 Ile Leu Asn Asp Asn Leu Ile Ser Phe Leu Pro Asp Asn Ile Phe Arg
 180 185 190
 Phe Ala Ser Leu Thr His Leu Asp Ile Arg Gly Asn Arg Ile Gln Lys
 195 200 205
 Leu Pro Tyr Ile Gly Val Leu Glu His Ile Gly Arg Val Val Glu Leu
 210 215 220
 Gln Leu Glu Asp Asn Pro Trp Asn Cys Ser Cys Asp Leu Leu Pro Leu
 225 230 235 240
 Lys Ala Trp Leu Glu Asn Met Pro Tyr Asn Ile Tyr Ile Gly Glu Ala
 245 250 255
 Ile Cys Glu Thr Pro Ser Asp Leu Tyr Gly Arg Leu Leu Lys Glu Thr
 260 265 270
 Asn Lys Gln Glu Leu Cys Pro Met Gly Thr Gly Ser Asp Phe Asp Val
 275 280 285
 Arg Ile Leu Pro Pro Ser Gln Leu Glu Asn Gly Tyr Thr Thr Pro Asn
 290 295 300
 Gly His Thr Thr Gln Thr Ser Leu His Arg Leu Val Thr Lys Pro Pro
 305 310 315 320
 Lys Thr Thr Asn Pro Ser Lys Ile Ser Gly Ile Val Ala Gly Lys Ala
 325 330 335
 Leu Ser Asn Arg Asn Leu Ser Gln Ile Val Ser Tyr Gln Thr Arg Val
 340 345 350
 Pro Pro Leu Thr Pro Cys Pro Ala Pro Cys Phe Cys Lys Thr His Pro
 355 360 365
 Ser Asp Leu Gly Leu Ser Val Asn Cys Gln Glu Lys Asn Ile Gln Ser
 370 375 380
 Met Ser Glu Leu Ile Pro Lys Pro Leu Asn Ala Lys Lys Leu His Val
 385 390 395 400
 Asn Gly Asn Ser Ile Lys Asp Val Asp Val Ser Asp Phe Thr Asp Phe
 405 410 415

Glu Gly Leu Asp Leu Leu His Leu Gly Ser Asn Gln Ile Thr Val Ile
 420 425 430
 Lys Gly Asp Val Phe His Asn Leu Thr Asn Leu Arg Arg Leu Tyr Leu
 435 440 445
 Asn Gly Asn Gln Ile Glu Arg Leu Tyr Pro Glu Ile Phe Ser Gly Leu
 450 455 460
 His Asn Leu Gln Tyr Leu Tyr Leu Glu Tyr Asn Leu Ile Lys Glu Ile
 465 470 475 480
 Ser Ala Gly Thr Phe Asp Ser Met Pro Asn Leu Gln Leu Leu Tyr Leu
 485 490 495
 Asn Asn Asn Leu Leu Lys Ser Leu Pro Val Tyr Ile Phe Ser Gly Ala
 500 505 510
 Pro Leu Ala Arg Leu Asn Leu Arg Asn Asn Lys Phe Met Tyr Leu Pro
 515 520 525
 Val Ser Gly Val Leu Asp Gln Leu Gln Ser Leu Thr Gln Ile Asp Leu
 530 535 540
 Glu Gly Asn Pro Trp Asp Cys Thr Cys Asp Leu Val Ala Leu Lys Leu
 545 550 555 560
 Trp Val Glu Lys Leu Ser Asp Gly Ile Val Val Lys Glu Leu Lys Cys
 565 570 575
 Glu Thr Pro Val Gln Phe Ala Asn Ile Glu Leu Lys Ser Leu Lys Asn
 580 585 590
 Glu Ile Leu Cys Pro Lys Leu Leu Asn Lys Pro Ser Ala Pro Phe Thr
 595 600 605
 Ser Pro Ala Pro Ala Ile Thr Phe Thr Thr Pro Leu Gly Pro Ile Arg
 610 615 620
 Ser Pro Pro Gly Gly Pro Val Pro Leu Ser Ile Leu Ile Leu Ser Ile
 625 630 635 640
 Leu Val Val Leu Ile Leu Thr Val Phe Val Ala Phe Cys Leu Leu Val
 645 650 655
 Phe Val Leu Arg Arg Asn Lys Lys Pro Thr Val Lys His Glu Gly Leu
 660 665 670
 Gly Asn Pro Asp Cys Gly Ser Met Gln Leu Gln Leu Arg Lys His Asp
 675 680 685
 His Lys Thr Asn Lys Lys Asp Gly Leu Ser Thr Glu Ala Phe Ile Pro
 690 695 700
 Gln Thr Ile Glu Gln Met Ser Lys Ser His Thr Cys Gly Leu Lys Glu
 705 710 715 720

Ser Glu Thr Gly Phe Met Phe Ser Asp Pro Pro Gly Gln Lys Val Val
 725 730 735
 Met Arg Asn Val Ala Asp Lys Glu Lys Asp Leu Leu His Val Asp Thr
 740 745 750
 Arg Lys Arg Leu Ser Thr Ile Asp Glu Leu Asp Glu Leu Phe Pro Ser
 755 760 765
 Arg Asp Ser Asn Val Phe Ile Gln Asn Phe Leu Glu Ser Lys Lys Glu
 770 775 780
 Tyr Asn Ser Ile Gly Val Ser Gly Phe Glu Ile Arg Tyr Pro Glu Lys
 785 790 795 800
 Gln Pro Asp Lys Lys Ser Lys Lys Ser Leu Ile Gly Gly Asn His Ser
 805 810 815
 Lys Ile Val Val Glu Gln Arg Lys Ser Glu Tyr Phe Glu Leu Lys Ala
 820 825 830
 Lys Leu Gln Ser Ser Pro Asp Tyr Leu Gln Val Leu Glu Glu Gln Thr
 835 840 845
 Ala Leu Asn Lys Ile
 850

<210> 31
 <211> 314
 <212> PRT
 <213> Homo sapiens

<400> 31
 Leu Gln Gln Trp Ile Gln Lys Leu Ser Lys Asn Thr Val Thr Asp Asp
 1 5 10 15
 Ile Leu Cys Thr Ser Pro Gly His Leu Asp Lys Lys Glu Leu Lys Ala
 20 25 30
 Leu Asn Ser Glu Ile Leu Cys Pro Gly Leu Val Asn Asn Pro Ser Met
 35 40 45
 Pro Thr Gln Thr Ser Tyr Leu Met Val Thr Thr Pro Ala Thr Thr Thr
 50 55 60
 Asn Thr Ala Asp Thr Ile Leu Arg Ser Leu Thr Asp Ala Val Pro Leu
 65 70 75 80
 Ser Val Leu Ile Leu Gly Leu Leu Ile Met Phe Ile Thr Ile Val Phe
 85 90 95
 Cys Ala Ala Gly Ile Val Val Leu Val Leu His Arg Arg Arg Arg Tyr
 100 105 110
 Lys Lys Lys Gln Val Asp Glu Gln Met Arg Asp Asn Ser Pro Val His
 115 120 125

Leu Gln Tyr Ser Met Tyr Gly His Lys Thr Thr His His Thr Thr Glu
 130 135 140
 Arg Pro Ser Ala Ser Leu Tyr Glu Gln His Met Val Ser Pro Met Val
 145 150 155 160
 His Val Tyr Arg Ser Pro Ser Phe Gly Pro Lys His Leu Glu Glu Glu
 165 170 175
 Glu Glu Arg Asn Glu Lys Glu Gly Ser Asp Ala Lys His Leu Gln Arg
 180 185 190
 Ser Leu Leu Glu Gln Glu Asn His Ser Pro Leu Thr Gly Ser Asn Met
 195 200 205
 Lys Tyr Lys Thr Thr Asn Gln Ser Thr Glu Phe Leu Ser Phe Gln Asp
 210 215 220
 Ala Ser Ser Leu Tyr Arg Asn Ile Leu Glu Lys Glu Arg Glu Leu Gln
 225 230 235 240
 Gln Leu Gly Ile Thr Glu Tyr Leu Arg Lys Asn Ile Ala Gln Leu Gln
 245 250 255
 Pro Asp Met Glu Ala His Tyr Pro Gly Ala His Glu Glu Leu Lys Leu
 260 265 270
 Met Glu Thr Leu Met Tyr Ser Arg Pro Arg Lys Val Leu Val Glu Gln
 275 280 285
 Thr Lys Asn Glu Tyr Phe Glu Leu Lys Ala Asn Leu His Ala Glu Pro
 290 295 300
 Asp Tyr Leu Glu Val Leu Glu Gln Gln Thr
 305 310

<210> 32
 <211> 966
 <212> PRT
 <213> Homo sapiens

<400> 32
 Arg Arg Gly Ala Gln Gly Gly Lys Met His Thr Cys Cys Pro Pro Val
 1 5 10 15
 Thr Leu Glu Gln Asp Leu His Arg Lys Met His Ser Trp Met Leu Gln
 20 25 30
 Thr Leu Ala Phe Ala Val Thr Ser Leu Val Leu Ser Cys Ala Glu Thr
 35 40 45
 Ile Asp Tyr Tyr Gly Glu Ile Cys Asp Asn Ala Cys Pro Cys Glu Glu
 50 55 60
 Lys Asp Gly Ile Leu Thr Val Ser Cys Glu Asn Arg Gly Ile Ile Ser

65	70	75	80
Leu Ser Glu Ile Ser Pro Pro Arg Phe Pro Ile Tyr His Leu Leu Leu	85	90	95
Ser Gly Asn Leu Leu Asn Arg Leu Tyr Pro Asn Glu Phe Val Asn Tyr	100	105	110
Thr Gly Ala Ser Ile Leu His Leu Gly Ser Asn Val Ile Gln Asp Ile	115	120	125
Glu Thr Gly Ala Phe His Gly Leu Arg Gly Leu Arg Arg Leu His Leu	130	135	140
Asn Asn Asn Lys Leu Glu Leu Leu Arg Asp Asp Thr Phe Leu Gly Leu	145	150	155
Glu Asn Leu Glu Tyr Leu Gln Val Asp Tyr Asn Tyr Ile Ser Val Ile	165	170	175
Glu Pro Asn Ala Phe Gly Lys Leu His Leu Leu Gln Val Leu Ile Leu	180	185	190
Asn Asp Asn Leu Leu Ser Ser Leu Pro Asn Asn Leu Phe Arg Phe Val	195	200	205
Pro Leu Thr His Leu Asp Leu Arg Gly Asn Arg Leu Lys Leu Leu Pro	210	215	220
Tyr Val Gly Leu Leu Gln His Met Asp Lys Val Val Glu Leu Gln Leu	225	230	235
Glu Glu Asn Pro Trp Asn Cys Ser Cys Glu Leu Ile Ser Leu Lys Asp	245	250	255
Trp Leu Asp Ser Ile Ser Tyr Ser Ala Leu Val Gly Asp Val Val Cys	260	265	270
Glu Thr Pro Phe Arg Leu His Gly Arg Asp Leu Asp Glu Val Ser Lys	275	280	285
Gln Glu Leu Cys Pro Arg Arg Leu Ile Ser Asp Tyr Glu Met Arg Pro	290	295	300
Gln Thr Pro Leu Ser Thr Thr Gly Tyr Leu His Thr Thr Pro Ala Ser	305	310	315
Val Asn Ser Val Ala Thr Ser Ser Ser Ala Val Tyr Lys Pro Pro Leu	325	330	335
Lys Pro Pro Lys Gly Thr Arg Gln Pro Asn Lys Pro Arg Val Arg Pro	340	345	350
Thr Ser Arg Gln Pro Ser Lys Asp Leu Gly Tyr Ser Asn Tyr Gly Pro	355	360	365
Ser Ile Ala Tyr Gln Thr Lys Ser Pro Val Pro Leu Glu Cys Pro Thr			

370					375					380					
Ala	Cys	Ser	Cys	Asn	Leu	Gln	Ile	Ser	Asp	Leu	Gly	Leu	Asn	Val	Asn
385					390					395					400
Cys	Gln	Glu	Arg	Lys	Ile	Glu	Ser	Ile	Ala	Glu	Leu	Gln	Pro	Lys	Pro
				405					410					415	
Tyr	Asn	Pro	Lys	Lys	Met	Tyr	Leu	Thr	Glu	Asn	Tyr	Ile	Ala	Val	Val
			420					425					430		
Arg	Arg	Thr	Asp	Phe	Leu	Glu	Ala	Thr	Gly	Leu	Asp	Leu	Leu	His	Leu
		435					440					445			
Gly	Asn	Asn	Arg	Ile	Ser	Met	Ile	Gln	Asp	Arg	Ala	Phe	Gly	Asp	Leu
	450					455					460				
Thr	Asn	Leu	Arg	Arg	Leu	Tyr	Leu	Asn	Gly	Asn	Arg	Ile	Glu	Arg	Leu
465					470					475					480
Ser	Pro	Glu	Leu	Phe	Tyr	Gly	Leu	Gln	Ser	Leu	Gln	Tyr	Leu	Phe	Leu
				485					490					495	
Gln	Tyr	Asn	Leu	Ile	Arg	Glu	Ile	Gln	Ser	Gly	Thr	Phe	Asp	Pro	Val
			500					505					510		
Pro	Asn	Leu	Gln	Leu	Leu	Phe	Leu	Asn	Asn	Asn	Leu	Leu	Gln	Ala	Met
		515					520					525			
Pro	Ser	Gly	Val	Phe	Ser	Gly	Leu	Thr	Leu	Leu	Arg	Leu	Asn	Leu	Arg
	530					535					540				
Ser	Asn	His	Phe	Thr	Ser	Leu	Pro	Val	Ser	Gly	Val	Leu	Asp	Gln	Leu
545					550					555					560
Lys	Ser	Leu	Ile	Gln	Ile	Asp	Leu	His	Asp	Asn	Pro	Trp	Asp	Cys	Thr
			565						570					575	
Cys	Asp	Ile	Val	Gly	Met	Lys	Leu	Trp	Val	Glu	Gln	Leu	Lys	Val	Gly
			580					585					590		
Val	Leu	Val	Asp	Glu	Val	Ile	Cys	Lys	Ala	Pro	Lys	Lys	Phe	Ala	Glu
		595					600					605			
Thr	Asp	Met	Arg	Ser	Ile	Lys	Ser	Glu	Leu	Leu	Cys	Pro	Asp	Tyr	Ser
	610					615					620				
Asp	Val	Val	Val	Ser	Thr	Pro	Thr	Pro	Ser	Ser	Ile	Gln	Val	Pro	Ala
625					630					635					640
Arg	Thr	Ser	Ala	Val	Thr	Pro	Ala	Val	Arg	Leu	Asn	Ser	Thr	Gly	Ala
			645						650					655	
Pro	Ala	Ser	Leu	Gly	Ala	Gly	Gly	Gly	Ala	Ser	Ser	Val	Pro	Leu	Ser
			660				665						670		
Val	Leu	Ile	Leu	Ser	Leu	Leu	Leu	Val	Phe	Ile	Met	Ser	Val	Phe	Val

675					680					685						
Ala	Ala	Gly	Leu	Phe	Val	Leu	Val	Met	Lys	Arg	Arg	Lys	Lys	Asn	Gln	
690					695					700						
Ser	Asp	His	Thr	Ser	Thr	Asn	Asn	Ser	Asp	Val	Ser	Ser	Phe	Asn	Met	
705					710					715					720	
Gln	Tyr	Ser	Val	Tyr	Gly	Gly	Gly	Gly	Gly	Thr	Gly	Gly	His	Pro	His	
725					730					735						
Ala	His	Val	His	His	Arg	Gly	Pro	Ala	Leu	Pro	Lys	Val	Lys	Thr	Pro	
740					745					750						
Ala	Gly	His	Val	Tyr	Glu	Tyr	Ile	Pro	His	Pro	Leu	Gly	His	Met	Cys	
755					760					765						
Lys	Asn	Pro	Ile	Tyr	Arg	Ser	Arg	Glu	Gly	Asn	Ser	Val	Glu	Asp	Tyr	
770					775					780						
Lys	Asp	Leu	His	Glu	Leu	Lys	Val	Thr	Tyr	Ser	Ser	Asn	His	His	Leu	
785					790					795					800	
Gln	Gln	Gln	Gln	Gln	Pro	Pro	Pro	Pro	Pro	Gln	Gln	Pro	Gln	Gln	Gln	
805					810					815						
Pro	Pro	Pro	Gln	Leu	Gln	Leu	Gln	Pro	Gly	Glu	Glu	Glu	Arg	Arg	Glu	
820					825					830						
Ser	His	His	Leu	Arg	Ser	Pro	Ala	Tyr	Ser	Val	Ser	Thr	Ile	Glu	Pro	
835					840					845						
Arg	Glu	Asp	Leu	Leu	Ser	Pro	Val	Gln	Asp	Ala	Asp	Arg	Phe	Tyr	Arg	
850					855					860						
Gly	Ile	Leu	Glu	Pro	Asp	Lys	His	Cys	Ser	Thr	Thr	Pro	Ala	Gly	Asn	
865					870					875					880	
Ser	Leu	Pro	Glu	Tyr	Pro	Lys	Phe	Pro	Cys	Ser	Pro	Ala	Ala	Tyr	Thr	
885					890					895						
Phe	Ser	Pro	Asn	Tyr	Asp	Leu	Arg	Arg	Pro	His	Gln	Tyr	Leu	His	Pro	
900					905					910						
Gly	Ala	Gly	Asp	Ser	Arg	Leu	Arg	Glu	Pro	Val	Leu	Tyr	Ser	Pro	Pro	
915					920					925						
Ser	Ala	Val	Phe	Val	Glu	Pro	Asn	Arg	Asn	Glu	Tyr	Leu	Glu	Leu	Lys	
930					935					940						
Ala	Lys	Leu	Asn	Val	Glu	Pro	Asp	Tyr	Leu	Glu	Val	Leu	Glu	Lys	Gln	
945					950					955					960	
Thr	Thr	Phe	Ser	Gln	Phe											
965																

<210> 33
 <211> 1428
 <212> PRT
 <213> Mus musculus

<400> 33

```

Met Val Ala Val Ala Val Ala Ala Ser Thr Glu Ala Arg Leu Arg Gly
  1              5              10              15

Ser Thr Thr Ala Thr Ala Ala Pro Ala Gly Arg Lys Gly Arg Gln His
      20              25              30

Arg Pro Cys Thr Ala Thr Gly Ala Trp Arg Pro Gly Pro Arg Ala Arg
      35              40              45

Leu Cys Leu Pro Arg Val Leu Ser Arg Ala Leu Pro Pro Pro Pro Leu
      50              55              60

Leu Pro Leu Leu Phe Ser Leu Leu Leu Leu Pro Leu Pro Arg Glu Ala
      65              70              75              80

Glu Ala Ala Ala Val Ala Ala Ala Val Ser Gly Ser Ala Ala Ala Glu
      85              90              95

Ala Lys Glu Cys Asp Arg Pro Cys Val Asn Gly Gly Arg Cys Asn Pro
      100             105             110

Gly Thr Gly Gln Cys Val Cys Pro Thr Gly Trp Val Gly Glu Gln Cys
      115             120             125

Gln His Cys Gly Gly Arg Phe Arg Leu Thr Gly Ser Ser Gly Phe Val
      130             135             140

Thr Asp Gly Pro Gly Asn Tyr Lys Tyr Lys Thr Lys Cys Thr Trp Leu
      145             150             155             160

Ile Glu Gly Tyr Pro Asn Ala Val Leu Arg Leu Arg Phe Asn His Phe
      165             170             175

Ala Thr Glu Cys Ser Trp Asp His Leu Tyr Val Tyr Asp Gly Asp Ser
      180             185             190

Ile Tyr Ala Pro Leu Ile Ala Ala Phe Ser Gly Leu Ile Val Pro Glu
      195             200             205

Arg Asp Gly Asn Glu Thr Ala Pro Glu Val Thr Val Thr Ser Gly Tyr
      210             215             220

Ala Leu Leu His Phe Phe Ser Asp Ala Ala Tyr Asn Leu Thr Gly Phe
      225             230             235             240

Asn Ile Thr Tyr Asn Phe Asp Met Cys Pro Asn Asn Cys Ser Ala Arg
      245             250             255

Gly Glu Cys Lys Ser Ser Asn Ser Ser Ser Ala Val Glu Cys Glu Cys
      260             265             270

```

Ser Glu Asn Trp Lys Gly Glu Ser Cys Asp Ile Pro His Cys Thr Asp
 275 280 285
 Asn Cys Gly Phe Pro His Arg Gly Ile Cys Asn Ala Ser Asp Thr Arg
 290 295 300
 Gly Cys Ser Cys Phe Pro His Trp Gln Gly Pro Gly Cys Ser Ile Pro
 305 310 315 320
 Val Pro Ala Asn Gln Ser Phe Trp Thr Arg Glu Glu Tyr Ser Asp Leu
 325 330 335
 Lys Leu Pro Arg Ala Ser His Lys Ala Val Val Asn Gly Asn Ile Met
 340 345 350
 Trp Val Val Gly Gly Tyr Met Phe Asn His Ser Asp Tyr Ser Met Val
 355 360 365
 Leu Ala Tyr Asp Leu Thr Ser Arg Glu Trp Leu Pro Leu Asn His Ser
 370 375 380
 Val Asn Ser Val Val Val Arg Tyr Gly His Ser Leu Ala Leu His Lys
 385 390 395 400
 Asp Lys Ile Tyr Met Tyr Gly Gly Lys Ile Asp Ser Thr Gly Asn Val
 405 410 415
 Thr Asn Glu Leu Arg Val Phe His Ile His Asn Glu Ser Trp Val Leu
 420 425 430
 Leu Thr Pro Lys Ala Lys Asp Gln Tyr Ala Val Val Gly His Ser Ala
 435 440 445
 His Ile Val Thr Leu Ala Ser Gly Arg Val Val Met Leu Val Ile Phe
 450 455 460
 Gly His Cys Pro Leu Tyr Gly Tyr Ile Ser Val Val Gln Glu Tyr Asp
 465 470 475 480
 Leu Glu Lys Asn Thr Trp Ser Ile Leu His Thr Gln Gly Ala Leu Val
 485 490 495
 Gln Gly Gly Tyr Gly His Ser Ser Ala Tyr Asp Asp Arg Thr Lys Ala
 500 505 510
 Leu Tyr Val His Gly Gly Tyr Lys Ala Phe Ser Ala Asn Lys Tyr Arg
 515 520 525
 Leu Ala Asp Asp Leu Tyr Arg Tyr Asp Val Asp Thr Gln Met Trp Thr
 530 535 540
 Ile Leu Lys Asp Ser Arg Phe Phe Arg Tyr Leu His Thr Ala Val Ile
 545 550 555 560
 Val Ser Gly Thr Met Leu Val Phe Gly Gly Asn Thr His Asn Asp Thr
 565 570 575

Ser Met Ser His Gly Ala Lys Cys Phe Ser Ser Asp Phe Met Ala Tyr
 580 585 590
 Asp Ile Ala Cys Asp Arg Trp Ser Val Leu Pro Arg Pro Glu Leu His
 595 600 605
 His Asp Val Asn Arg Phe Gly His Ser Ala Val Leu Tyr Asn Ser Thr
 610 615 620
 Met Tyr Val Phe Gly Gly Phe Asn Ser Leu Leu Leu Ser Asp Val Leu
 625 630 635 640
 Val Phe Thr Ser Glu Gln Cys Asp Ala His Arg Ser Glu Ala Ala Cys
 645 650 655
 Val Ala Ala Gly Pro Gly Ile Arg Cys Leu Trp Asp Thr Gln Ser Ser
 660 665 670
 Arg Cys Thr Ser Trp Glu Leu Ala Thr Glu Glu Gln Ala Glu Lys Leu
 675 680 685
 Lys Ser Glu Cys Phe Ser Lys Arg Thr Leu Asp His Asp Arg Cys Asp
 690 695 700
 Gln His Thr Asp Cys Tyr Ser Cys Thr Ala Asn Thr Asn Asp Cys His
 705 710 715 720
 Trp Cys Asn Asp His Cys Val Pro Val Asn His Ser Cys Thr Glu Gly
 725 730 735
 Gln Ile Ser Ile Ala Lys Tyr Glu Ser Cys Pro Lys Asp Asn Pro Met
 740 745 750
 Tyr Tyr Cys Asn Lys Lys Thr Ser Cys Arg Ser Cys Ala Leu Asp Gln
 755 760 765
 Asn Cys Gln Trp Glu Pro Arg Asn Gln Glu Cys Ile Ala Leu Pro Glu
 770 775 780
 Asn Ile Cys Gly Asn Gly Trp His Leu Val Gly Asn Ser Cys Leu Lys
 785 790 795 800
 Ile Thr Thr Ala Lys Glu Asn Tyr Asp Asn Ala Lys Leu Ser Cys Arg
 805 810 815
 Asn His Asn Ala Phe Leu Ala Ser Leu Thr Ser Gln Lys Lys Val Glu
 820 825 830
 Leu Val Leu Lys Gln Leu Arg Leu Met Gln Ser Ser Gln Ser Met Ser
 835 840 845
 Lys Leu Thr Leu Thr Pro Trp Val Gly Leu Arg Lys Ile Asn Val Ser
 850 855 860
 Tyr Trp Cys Trp Glu Asp Met Ser Pro Phe Thr Asn Ser Leu Leu Gln
 865 870 875 880

Trp Met Pro Ser Glu Pro Ser Asp Ala Gly Phe Cys Gly Ile Leu Ser
 885 890 895

Glu Pro Ser Thr Arg Gly Leu Lys Ala Ala Thr Cys Ile Asn Pro Leu
 900 905 910

Asn Gly Ser Val Cys Glu Arg Pro Ala Asn His Ser Ala Lys Gln Cys
 915 920 925

Arg Thr Pro Cys Ala Leu Arg Thr Ala Cys Gly Glu Cys Thr Ser Ser
 930 935 940

Ser Ser Glu Cys Met Trp Cys Ser Asn Met Lys Gln Cys Val Asp Ser
 945 950 955 960

Asn Ala Tyr Val Ala Ser Phe Pro Phe Gly Gln Cys Met Glu Trp Tyr
 965 970 975

Thr Met Ser Ser Cys Pro Pro Glu Asn Cys Ser Gly Tyr Cys Thr Cys
 980 985 990

Ser His Cys Leu Glu Gln Pro Gly Cys Gly Trp Cys Thr Asp Pro Ser
 995 1000 1005

Asn Thr Gly Lys Gly Lys Cys Ile Glu Gly Ser Tyr Lys Gly Pro Val
 1010 1015 1020

Lys Met Pro Ser Gln Ala Ser Ala Gly Asn Val Tyr Pro Gln Pro Leu
 1025 1030 1035 1040

Leu Asn Ser Ser Met Cys Leu Glu Asp Ser Arg Tyr Asn Trp Ser Phe
 1045 1050 1055

Ile His Cys Pro Ala Cys Gln Cys Asn Gly His Ser Lys Cys Ile Asn
 1060 1065 1070

Gln Ser Val Cys Glu Lys Cys Glu Asp Leu Thr Thr Gly Lys His Cys
 1075 1080 1085

Glu Thr Cys Ile Ser Gly Phe Tyr Gly Asp Pro Thr Asn Gly Gly Lys
 1090 1095 1100

Cys Gln Pro Cys Lys Cys Asn Gly His Ala Ser Leu Cys Asn Thr Asn
 1105 1110 1115 1120

Thr Gly Lys Cys Phe Cys Thr Thr Lys Gly Val Lys Gly Asp Glu Cys
 1125 1130 1135

Gln Leu Cys Lys Val Glu Asn Arg Tyr Gln Gly Asn Pro Leu Lys Gly
 1140 1145 1150

Thr Cys Tyr Tyr Thr Leu Leu Ile Asp Tyr Gln Phe Thr Phe Ser Leu
 1155 1160 1165

Ser Gln Gly Asp Asp Arg Tyr Tyr Thr Ala Ile Asn Phe Val Ala Thr
 1170 1175 1180

Pro Asp Glu Gln Asn Arg Asp Phe Asp Met Phe Ile Asn Ala Ser Lys
 1185 1190 1195 1200
 Lys Phe Asn Leu Asn Ile Thr Trp Ala Thr Ser Phe Pro Ala Gly Thr
 1205 1210 1215
 Gln Thr Gly Glu Glu Val Pro Val Val Ser Lys Thr Asn Ile Lys Glu
 1220 1225 1230
 Tyr Lys Asp Ser Phe Ser Asn Glu Lys Phe Asp Phe Arg Asn His Pro
 1235 1240 1245
 Asn Ile Thr Phe Phe Val Tyr Val Ser Asn Phe Thr Trp Pro Ile Lys
 1250 1255 1260
 Ile Gln Ile Ala Phe Ser Gln His Ser Asn Phe Met Asp Leu Val Gln
 1265 1270 1275 1280
 Phe Phe Val Thr Phe Phe Ser Cys Phe Leu Ser Leu Leu Leu Val Ala
 1285 1290 1295
 Ala Val Val Trp Lys Ile Lys Gln Ser Cys Trp Ala Ser Arg Arg Arg
 1300 1305 1310
 Glu Gln Leu Leu Arg Glu Met Gln Gln Met Ala Ser Arg Pro Phe Ala
 1315 1320 1325
 Ser Val Asn Val Ala Leu Glu Thr Asp Glu Glu Pro Pro Asp Leu Ile
 1330 1335 1340
 Gly Gly Ser Ile Lys Thr Val Pro Lys Pro Ile Ala Leu Glu Pro Cys
 1345 1350 1355 1360
 Phe Gly Asn Lys Ala Ala Val Leu Ser Val Phe Val Arg Leu Pro Arg
 1365 1370 1375
 Gly Leu Gly Gly Ile Pro Pro Pro Gly Gln Ser Gly Leu Ala Val Ala
 1380 1385 1390
 Ser Ala Leu Val Asp Ile Ser Gln Gln Met Pro Ile Val Tyr Lys Glu
 1395 1400 1405
 Lys Ser Gly Ala Val Arg Asn Arg Lys Gln Gln Pro Pro Ala Gln Pro
 1410 1415 1420
 Gly Thr Cys Ile
 1425

<210> 34

<211> 1428

<212> PRT

<213> Mus musculus

<400> 34

Met Val Ala Val Ala Ala Ala Ala Ala Thr Glu Ala Arg Leu Arg Gly
 1 5 10 15

Ser Thr Thr Thr Thr Ala Ala Pro Ala Gly Arg Lys Gly Arg Gln His
 20 25 30
 Arg Pro Cys Thr Ala Thr Gly Ala Trp Arg Pro Gly Pro Arg Ala Arg
 35 40 45
 Leu Cys Leu Pro Arg Val Leu Ser Arg Ala Leu Pro Pro Pro Pro Leu
 50 55 60
 Leu Pro Leu Leu Phe Ser Leu Leu Leu Leu Pro Leu Pro Arg Glu Ala
 65 70 75 80
 Glu Ala Ala Ala Val Ala Ala Ala Val Ser Gly Ser Ala Ala Ala Glu
 85 90 95
 Ala Lys Glu Cys Asp Arg Pro Cys Val Asn Gly Gly Arg Cys Asn Pro
 100 105 110
 Gly Thr Gly Gln Cys Val Cys Pro Thr Gly Trp Val Gly Glu Gln Cys
 115 120 125
 Gln His Cys Gly Gly Arg Phe Arg Leu Thr Gly Ser Ser Gly Phe Val
 130 135 140
 Thr Asp Gly Pro Gly Asn Tyr Lys Tyr Lys Thr Lys Cys Thr Trp Leu
 145 150 155 160
 Ile Glu Gly Gln Pro Asn Arg Ile Met Arg Leu Arg Phe Asn His Phe
 165 170 175
 Ala Thr Glu Cys Ser Trp Asp His Leu Tyr Val Tyr Asp Gly Asp Ser
 180 185 190
 Ile Tyr Ala Pro Leu Ile Ala Ala Phe Ser Gly Leu Ile Val Pro Glu
 195 200 205
 Arg Asp Gly Asn Glu Thr Ala Pro Glu Val Thr Val Thr Ser Gly Tyr
 210 215 220
 Ala Leu Leu His Phe Phe Ser Asp Ala Ala Tyr Asn Leu Thr Gly Phe
 225 230 235 240
 Asn Ile Thr Tyr Asn Phe Asp Met Cys Pro Asn Asn Cys Ser Gly Arg
 245 250 255
 Gly Glu Cys Lys Ser Ser Asn Ser Ser Ser Ala Val Glu Cys Glu Cys
 260 265 270
 Ser Glu Asn Trp Lys Gly Glu Ser Cys Asp Ile Pro His Cys Thr Asp
 275 280 285
 Asn Cys Gly Phe Pro His Arg Gly Ile Cys Asn Ala Ser Asp Thr Arg
 290 295 300
 Gly Cys Ser Cys Phe Pro His Trp Gln Gly Pro Gly Cys Ser Ile Pro
 305 310 315 320

Val Pro Ala Asn Gln Ser Phe Trp Thr Arg Glu Glu Tyr Ser Asp Leu
 325 330 335
 Lys Leu Pro Arg Ala Ser His Lys Ala Val Val Asn Gly Asn Ile Met
 340 345 350
 Trp Val Val Gly Gly Tyr Met Phe Asn His Ser Asp Tyr Ser Met Val
 355 360 365
 Leu Ala Tyr Asp Leu Thr Ser Arg Glu Trp Leu Pro Leu Asn His Ser
 370 375 380
 Val Asn Ser Val Val Val Arg Tyr Gly His Ser Leu Ala Leu His Lys
 385 390 395 400
 Asp Lys Ile Tyr Met Tyr Gly Gly Lys Ile Asp Ser Thr Gly Asn Val
 405 410 415
 Thr Asn Glu Leu Arg Val Phe His Ile His Asn Glu Ser Trp Val Leu
 420 425 430
 Leu Thr Pro Lys Ala Lys Asp Gln Tyr Ala Val Val Gly His Ser Ala
 435 440 445
 His Ile Val Thr Leu Ala Ser Gly Arg Val Val Met Leu Val Ile Phe
 450 455 460
 Gly His Cys Pro Leu Tyr Gly Tyr Ile Ser Val Val Gln Glu Tyr Asp
 465 470 475 480
 Leu Glu Lys Asn Thr Trp Ser Ile Leu His Thr Gln Gly Ala Leu Val
 485 490 495
 Gln Gly Gly Tyr Gly His Ser Ser Val Tyr Asp Asp Arg Thr Lys Ala
 500 505 510
 Leu Tyr Val His Gly Gly Tyr Lys Ala Phe Ser Ala Asn Lys Tyr Arg
 515 520 525
 Leu Ala Asp Asp Leu Tyr Arg Tyr Asp Val Asp Thr Gln Met Trp Thr
 530 535 540
 Ile Leu Lys Asp Ser Arg Phe Phe Arg Tyr Leu His Thr Ala Val Ile
 545 550 555 560
 Val Ser Gly Thr Met Leu Val Phe Gly Gly Asn Thr His Asn Asp Thr
 565 570 575
 Ser Met Ser His Gly Ala Lys Cys Phe Ser Ser Asp Phe Met Ala Tyr
 580 585 590
 Asp Ile Ala Cys Asp Arg Trp Ser Val Leu Pro Arg Pro Glu Leu His
 595 600 605
 His Asp Val Asn Arg Phe Gly His Ser Ala Val Leu Tyr Asn Ser Thr
 610 615 620

Met	Tyr	Val	Phe	Gly	Gly	Phe	Asn	Ser	Leu	Leu	Leu	Ser	Asp	Val	Leu	625	630	635	640
Val	Phe	Thr	Ser	Glu	Gln	Cys	Asp	Ala	His	Arg	Ser	Glu	Ala	Ala	Cys	645	650	655	
Val	Ala	Ala	Gly	Pro	Gly	Ile	Arg	Cys	Leu	Trp	Asp	Thr	Gln	Ser	Ser	660	665	670	
Arg	Cys	Thr	Ser	Trp	Glu	Leu	Ala	Thr	Glu	Glu	Gln	Ala	Glu	Lys	Leu	675	680	685	
Lys	Ser	Glu	Cys	Phe	Ser	Lys	Arg	Thr	Leu	Asp	His	Asp	Arg	Cys	Asp	690	695	700	
Gln	His	Thr	Asp	Cys	Tyr	Ser	Cys	Thr	Ala	Asn	Thr	Asn	Asp	Cys	His	705	710	715	720
Trp	Cys	Asn	Asp	His	Cys	Val	Pro	Val	Asn	His	Ser	Cys	Thr	Glu	Gly	725	730	735	
Gln	Ile	Ser	Ile	Ala	Lys	Tyr	Glu	Ser	Cys	Pro	Lys	Asp	Asn	Pro	Met	740	745	750	
Tyr	Tyr	Cys	Asn	Lys	Lys	Thr	Ser	Cys	Arg	Ser	Cys	Ala	Leu	Asp	Gln	755	760	765	
Asn	Cys	Gln	Trp	Glu	Pro	Arg	Asn	Gln	Glu	Cys	Ile	Ala	Leu	Pro	Glu	770	775	780	
Asn	Ile	Cys	Gly	Asn	Gly	Trp	His	Leu	Val	Gly	Asn	Ser	Cys	Leu	Lys	785	790	795	800
Ile	Thr	Thr	Ala	Lys	Glu	Asn	Tyr	Asp	Asn	Ala	Lys	Leu	Ser	Cys	Arg	805	810	815	
Asn	His	Asn	Ala	Phe	Leu	Ala	Ser	Leu	Thr	Ser	Gln	Lys	Lys	Val	Glu	820	825	830	
Phe	Val	Leu	Lys	Gln	Leu	Arg	Leu	Met	Gln	Ser	Ser	Gln	Ser	Met	Ser	835	840	845	
Lys	Leu	Thr	Leu	Thr	Pro	Trp	Val	Gly	Leu	Arg	Lys	Ile	Asn	Val	Ser	850	855	860	
Tyr	Trp	Cys	Trp	Glu	Asp	Met	Ser	Pro	Phe	Thr	Asn	Ser	Leu	Leu	Gln	865	870	875	880
Trp	Met	Pro	Ser	Glu	Pro	Ser	Asp	Ala	Gly	Phe	Cys	Gly	Ile	Leu	Ser	885	890	895	
Glu	Pro	Ser	Thr	Arg	Gly	Leu	Lys	Ala	Ala	Thr	Cys	Ile	Asn	Pro	Leu	900	905	910	
Asn	Gly	Ser	Val	Cys	Glu	Arg	Pro	Ala	Asn	His	Ser	Ala	Lys	Gln	Cys	915	920	925	

Arg Thr Pro Cys Ala Leu Arg Thr Ala Cys Gly Glu Cys Thr Ser Ser
 930 935 940
 Ser Ser Glu Cys Met Trp Cys Ser Asn Met Lys Gln Cys Val Asp Ser
 945 950 955 960
 Asn Ala Tyr Val Ala Ser Phe Pro Phe Gly Gln Cys Met Glu Trp Tyr
 965 970 975
 Thr Met Ser Ser Cys Pro Pro Glu Asn Cys Ser Gly Tyr Cys Thr Cys
 980 985 990
 Ser His Cys Leu Glu Gln Pro Gly Cys Gly Trp Cys Thr Asp Pro Ser
 995 1000 1005
 Asn Thr Gly Lys Gly Lys Cys Ile Glu Gly Ser Tyr Lys Gly Pro Val
 1010 1015 1020
 Lys Met Pro Ser Gln Ala Ser Ala Gly Asn Val Tyr Pro Gln Pro Leu
 1025 1030 1035 1040
 Leu Asn Ser Ser Met Cys Leu Glu Asp Ser Arg Tyr Asn Trp Ser Phe
 1045 1050 1055
 Ile His Cys Pro Ala Cys Gln Cys Asn Gly His Ser Lys Cys Ile Asn
 1060 1065 1070
 Gln Ser Ile Cys Glu Lys Cys Glu Asp Leu Thr Thr Gly Lys His Cys
 1075 1080 1085
 Glu Thr Cys Ile Ser Gly Phe Tyr Gly Asp Pro Thr Asn Gly Gly Lys
 1090 1095 1100
 Cys Gln Pro Cys Lys Cys Asn Gly His Ala Ser Leu Cys Asn Thr Asn
 1105 1110 1115 1120
 Thr Gly Lys Cys Phe Cys Thr Thr Lys Gly Val Lys Gly Asp Glu Cys
 1125 1130 1135
 Gln Leu Cys Glu Val Glu Asn Arg Tyr Gln Gly Asn Pro Leu Lys Gly
 1140 1145 1150
 Thr Cys Tyr Tyr Thr Leu Leu Ile Asp Tyr Gln Phe Thr Phe Ser Leu
 1155 1160 1165
 Ser Gln Glu Asp Asp Arg Tyr Tyr Thr Ala Ile Asn Phe Val Ala Thr
 1170 1175 1180
 Pro Asp Glu Gln Asn Arg Asp Leu Asp Met Phe Ile Asn Ala Ser Lys
 1185 1190 1195 1200
 Asn Phe Asn Leu Asn Ile Thr Trp Ala Thr Ser Phe Pro Ala Gly Thr
 1205 1210 1215
 Gln Thr Gly Glu Glu Val Pro Val Val Ser Lys Thr Asn Ile Lys Glu
 1220 1225 1230

Tyr Lys Asp Ser Phe Ser Asn Glu Lys Phe Asp Phe Arg Asn His Pro
 1235 1240 1245
 Asn Ile Thr Phe Phe Val Tyr Val Ser Asn Phe Thr Trp Pro Ile Lys
 1250 1255 1260
 Ile Gln Ile Ala Phe Ser Gln His Ser Asn Phe Met Asp Leu Val Gln
 1265 1270 1275 1280
 Phe Phe Val Thr Phe Phe Ser Cys Phe Leu Ser Leu Leu Leu Val Ala
 1285 1290 1295
 Ala Val Val Trp Lys Ile Lys Gln Ser Cys Trp Ala Ser Arg Arg Arg
 1300 1305 1310
 Glu Gln Leu Leu Arg Glu Met Gln Gln Met Ala Ser Arg Pro Phe Ala
 1315 1320 1325
 Ser Val Asn Val Ala Leu Glu Thr Asp Glu Glu Pro Pro Asp Leu Ile
 1330 1335 1340
 Gly Gly Ser Ile Lys Thr Val Pro Lys Pro Ile Ala Leu Glu Pro Cys
 1345 1350 1355 1360
 Phe Gly Asn Lys Ala Ala Val Leu Ser Val Phe Val Arg Leu Pro Arg
 1365 1370 1375
 Gly Leu Gly Gly Ile Pro Pro Pro Gly Gln Ser Gly Leu Ala Val Ala
 1380 1385 1390
 Ser Ala Leu Val Asp Ile Ser Gln Gln Met Pro Ile Val Tyr Lys Glu
 1395 1400 1405
 Lys Ser Gly Ala Val Arg Asn Arg Lys Gln Gln Pro Pro Ala Gln Pro
 1410 1415 1420
 Gly Thr Cys Ile
 1425

<210> 35
 <211> 1198
 <212> PRT
 <213> Homo sapiens

<400> 35
 Met Val Ala Ala Ala Ala Thr Glu Ala Arg Leu Arg Arg Arg Thr
 1 5 10 15
 Ala Ala Thr Ala Ala Leu Ala Gly Arg Ser Gly Gly Pro His Cys Val
 20 25 30
 Asn Gly Gly Arg Cys Asn Pro Gly Thr Gly Gln Cys Val Cys Pro Ala
 35 40 45
 Gly Trp Val Gly Glu Gln Cys Gln His Cys Gly Gly Arg Phe Arg Leu

50					55					60					
Thr	Gly	Ser	Ser	Gly	Phe	Val	Thr	Asp	Gly	Pro	Gly	Asn	Tyr	Lys	Tyr
65					70					75					80
Lys	Thr	Lys	Cys	Thr	Trp	Leu	Ile	Glu	Gly	Gln	Pro	Asn	Arg	Ile	Met
			85						90					95	
Arg	Leu	Arg	Phe	Asn	His	Phe	Ala	Thr	Glu	Cys	Ser	Trp	Asp	His	Leu
			100					105					110		
Tyr	Val	Tyr	Asp	Gly	Asp	Ser	Ile	Tyr	Ala	Pro	Leu	Val	Ala	Ala	Phe
		115					120					125			
Ser	Gly	Leu	Ile	Val	Pro	Glu	Arg	Asp	Gly	Asn	Glu	Thr	Val	Pro	Glu
	130					135					140				
Val	Val	Ala	Thr	Ser	Gly	Tyr	Ala	Leu	Leu	His	Phe	Phe	Ser	Asp	Ala
145					150					155					160
Ala	Tyr	Asn	Leu	Thr	Gly	Phe	Asn	Ile	Thr	Tyr	Ser	Phe	Asp	Met	Cys
			165					170						175	
Pro	Asn	Asn	Cys	Ser	Gly	Arg	Gly	Glu	Cys	Lys	Ile	Ser	Asn	Ser	Ser
			180					185					190		
Asp	Thr	Val	Glu	Cys	Glu	Cys	Ser	Glu	Asn	Trp	Lys	Gly	Glu	Ala	Cys
		195					200					205			
Asp	Ile	Pro	His	Cys	Thr	Asp	Asn	Cys	Gly	Phe	Pro	His	Arg	Gly	Ile
	210					215					220				
Cys	Asn	Ser	Ser	Asp	Val	Arg	Gly	Cys	Ser	Cys	Phe	Ser	Asp	Trp	Gln
225					230					235					240
Gly	Pro	Gly	Cys	Ser	Val	Pro	Val	Pro	Ala	Asn	Gln	Ser	Phe	Trp	Thr
			245						250					255	
Arg	Glu	Glu	Tyr	Ser	Asn	Leu	Lys	Leu	Pro	Arg	Ala	Ser	His	Lys	Ala
			260					265					270		
Val	Val	Asn	Gly	Asn	Ile	Met	Trp	Val	Val	Gly	Gly	Tyr	Met	Phe	Asn
		275					280					285			
His	Ser	Asp	Tyr	Asn	Met	Val	Leu	Ala	Tyr	Asp	Leu	Ala	Ser	Arg	Glu
	290					295					300				
Trp	Leu	Pro	Leu	Asn	Arg	Ser	Val	Asn	Asn	Val	Val	Val	Arg	Tyr	Gly
305					310					315					320
His	Ser	Leu	Ala	Leu	Tyr	Lys	Asp	Lys	Ile	Tyr	Met	Tyr	Gly	Gly	Lys
			325						330					335	
Ile	Asp	Ser	Thr	Gly	Asn	Val	Thr	Asn	Glu	Leu	Arg	Val	Phe	His	Ile
			340					345					350		
His	Asn	Glu	Ser	Trp	Val	Leu	Leu	Thr	Pro	Lys	Ala	Lys	Glu	Gln	Tyr

355	360	365
Ala Val Val Gly His Ser	Ala His Ile Val Thr	Leu Lys Asn Gly Arg
370	375	380
Val Val Met Leu Val	Ile Phe Gly His Cys	Pro Leu Tyr Gly Tyr Ile
385	390	395
Ser Asn Val Gln Glu Tyr	Asp Leu Asp Lys Asn	Thr Trp Ser Ile Leu
405	410	415
His Thr Gln Gly Ala Leu	Val Gln Gly Gly Tyr	Gly His Ser Ser Val
420	425	430
Tyr Asp His Arg Thr	Arg Ala Leu Tyr Val	His Gly Gly Tyr Lys Ala
435	440	445
Phe Ser Ala Asn Lys Tyr	Arg Leu Ala Asp Asp	Leu Tyr Arg Tyr Asp
450	455	460
Val Asp Thr Gln Met Trp	Thr Ile Leu Lys Asp	Ser Arg Phe Phe Arg
465	470	475
Tyr Leu His Thr Ala Val	Ile Val Ser Gly Thr	Met Leu Val Phe Gly
485	490	495
Gly Asn Thr His Asn Asp	Thr Ser Met Ser His	Gly Ala Lys Cys Phe
500	505	510
Ser Ser Asp Phe Met Ala	Tyr Asp Ile Ala Cys	Asp Arg Trp Ser Val
515	520	525
Leu Pro Arg Pro Asp Ser	Thr Met Met Ser Thr	Asp Leu Ala Ile Pro
530	535	540
Ala Val Leu His Asn Ser	Thr Met Tyr Val Phe	Gly Gly Phe Asn Ser
545	550	555
Leu Leu Leu Ser Asp Ile	Leu Val Phe Thr Ser	Glu Gln Cys Asp Ala
565	570	575
His Arg Ser Glu Ala Ala	Cys Leu Ala Ala Gly	Pro Gly Ile Arg Cys
580	585	590
Val Trp Asn Thr Gly Ser	Ser Ser Gln Cys Ile	Ser Trp Ala Leu Ala Thr
595	600	605
Asp Glu Gln Glu Glu Lys	Leu Lys Ser Glu Cys	Phe Ser Lys Arg Thr
610	615	620
Leu Asp His Asp Arg Cys	Asp Gln His Thr Asp	Cys Tyr Ser Cys Thr
625	630	635
Ala Asn Thr Asn Asp Cys	His Trp Cys Asn Asp	His Cys Val Pro Arg
645	650	655
Asn His Ser Cys Ser Glu	Gly Gln Ile Ser Ile	Phe Arg Tyr Glu Asn

660					665					670					
Cys	Pro	Lys	Asp	Asn	Pro	Met	Tyr	Tyr	Cys	Asn	Lys	Lys	Thr	Ser	Cys
675					680					685					
Arg	Ser	Cys	Ala	Leu	Asp	Gln	Asn	Cys	Gln	Trp	Glu	Pro	Arg	Asn	Gln
690					695					700					
Glu	Cys	Ile	Ala	Leu	Pro	Glu	Asn	Ile	Cys	Gly	Ile	Gly	Trp	His	Leu
705					710					715					
Val	Gly	Asn	Ser	Cys	Leu	Lys	Ile	Thr	Thr	Ala	Lys	Glu	Asn	Tyr	Asp
725					730					735					
Asn	Ala	Lys	Leu	Phe	Cys	Arg	Asn	His	Asn	Ala	Leu	Leu	Ala	Ser	Leu
740					745					750					
Thr	Thr	Gln	Lys	Lys	Val	Glu	Phe	Val	Leu	Lys	Gln	Leu	Arg	Ile	Met
755					760					765					
Gln	Ser	Ser	Gln	Ser	Met	Ser	Lys	Leu	Thr	Leu	Thr	Pro	Trp	Val	Gly
770					775					780					
Leu	Arg	Lys	Ile	Asn	Val	Ser	Tyr	Trp	Cys	Trp	Glu	Asp	Met	Ser	Pro
785					790					795					
Phe	Thr	Asn	Ser	Leu	Leu	Gln	Trp	Met	Pro	Ser	Glu	Pro	Ser	Asp	Ala
805					810					815					
Gly	Phe	Cys	Gly	Ile	Leu	Ser	Glu	Pro	Ser	Thr	Arg	Gly	Leu	Lys	Ala
820					825					830					
Ala	Thr	Cys	Ile	Asn	Pro	Leu	Asn	Gly	Ser	Val	Cys	Glu	Arg	Pro	Ala
835					840					845					
Asn	His	Ser	Ala	Lys	Gln	Cys	Arg	Thr	Pro	Cys	Ala	Leu	Arg	Thr	Ala
850					855					860					
Cys	Gly	Asp	Cys	Thr	Ser	Gly	Ser	Ser	Glu	Cys	Met	Trp	Cys	Ser	Asn
865					870					875					
Met	Lys	Gln	Cys	Val	Asp	Ser	Asn	Ala	Tyr	Val	Ala	Ser	Phe	Pro	Phe
885					890					895					
Gly	Gln	Cys	Met	Glu	Trp	Tyr	Thr	Met	Ser	Thr	Cys	Pro	Pro	Glu	Asn
900					905					910					
Cys	Ser	Gly	Tyr	Cys	Thr	Cys	Ser	His	Cys	Leu	Glu	Gln	Pro	Gly	Cys
915					920					925					
Gly	Trp	Cys	Thr	Asp	Pro	Ser	Asn	Thr	Gly	Lys	Gly	Lys	Cys	Ile	Glu
930					935					940					
Gly	Ser	Tyr	Lys	Gly	Pro	Val	Lys	Met	Pro	Ser	Gln	Ala	Pro	Thr	Gly
945					950					955					
Asn	Phe	Tyr	Pro	Gln	Pro	Leu	Leu	Asn	Ser	Ser	Met	Cys	Leu	Glu	Asp

His Arg Pro Cys Ala Ala Thr Gly Ala Trp Arg Pro Trp Pro Arg Ala
 35 40 45
 Gly Leu Cys Leu Pro Arg Val Leu Ser Arg Ala Leu Ser Pro Pro Pro
 50 55 60
 Leu Leu Pro Leu Leu Pro Leu Leu Phe Ser Leu Leu Leu Leu Pro Leu
 65 70 75 80
 Pro Arg Glu Ala Glu Ala Ala Ala Val Ala Ala Ala Val Ser Gly Ser
 85 90 95
 Ala Ala Ala Glu Ala Lys Glu Cys Asp Arg Pro Cys Val Asn Gly Gly
 100 105 110
 Arg Cys Asn Pro Gly Thr Gly Gln Cys Val Cys Pro Thr Gly Trp Val
 115 120 125
 Gly Glu Gln Cys Gln His Cys Gly Gly Arg Phe Arg Leu Thr Gly Ser
 130 135 140
 Ser Gly Phe Val Thr Asp Gly Pro Gly Asn Tyr Lys Tyr Lys Thr Lys
 145 150 155 160
 Cys Thr Trp Leu Ile Glu Gly Gln Pro Asn Lys Ile Met Arg Leu Arg
 165 170 175
 Phe Asn His Phe Ala Thr Glu Cys Ser Trp Asp His Leu Tyr Val Tyr
 180 185 190
 Asp Gly Asp Ser Ile Tyr Ala Pro Leu Ile Ala Ala Phe Ser Gly Leu
 195 200 205
 Ile Val Pro Glu Arg Asp Gly Asn Glu Thr Ala Pro Glu Val Thr Val
 210 215 220
 Thr Ser Gly Tyr Ala Leu Leu His Phe Phe Ser Asp Ala Ala Tyr Asn
 225 230 235 240
 Leu Thr Gly Phe Asn Ile Thr Tyr Asn Phe Asp Met Cys Pro Asn Asn
 245 250 255
 Cys Ser Gly Arg Gly Glu Cys Lys Ser Ser Asn Ser Ser Ser Thr Val
 260 265 270
 Glu Cys Glu Cys Ser Glu Asn Trp Lys Gly Glu Ser Cys Asp Ile Pro
 275 280 285
 His Cys Thr Asp Asn Cys Gly Phe Pro His Arg Gly Ile Cys Asn Ala
 290 295 300
 Ser Asp Thr Arg Gly Cys Ser Cys Phe Pro His Trp Gln Gly Pro Gly
 305 310 315 320
 Cys Ser Ile Pro Val Pro Ala Asn Gln Ser Phe Trp Thr Arg Glu Glu
 325 330 335

Tyr Ser Asp Leu Lys Leu Pro Arg Ala Ser His Lys Ala Glu Val Asn
 340 345 350
 Gly Asn Ile Met Trp Val Val Gly Gly Tyr Met Phe Asn His Ser Asp
 355 360 365
 Tyr Ser Met Val Leu Ala Tyr Asp Leu Ala Ser Arg Glu Trp Leu Ser
 370 375 380
 Leu Asn His Ser Val Asn Ser Val Val Val Arg Tyr Gly His Ser Leu
 385 390 395 400
 Ala Leu His Lys Asp Lys Ile Tyr Met Tyr Gly Gly Lys Ile Asp Ser
 405 410 415
 Thr Gly Asn Val Thr Asn Glu Leu Arg Val Phe His Ile His Asn Glu
 420 425 430
 Ser Trp Val Leu Leu Thr Pro Lys Ala Lys Asp Gln Tyr Ala Val Val
 435 440 445
 Gly His Ser Ala His Ile Val Thr Leu Ser Ser Gly Arg Val Val Met
 450 455 460
 Leu Val Ile Phe Gly His Cys Pro Leu Tyr Gly Tyr Ile Ser Val Val
 465 470 475 480
 Gln Glu Tyr Asp Leu Glu Lys Asn Thr Trp Ser Ile Leu Gln Thr Gln
 485 490 495
 Gly Ala Leu Val Gln Gly Gly Tyr Gly His Ser Ser Val Tyr Asp His
 500 505 510
 Arg Thr Lys Ala Leu Tyr Val His Gly Gly Tyr Lys Ala Phe Ser Ala
 515 520 525
 Asn Lys Tyr Arg Leu Ala Asp Asp Leu Tyr Arg Tyr His Val Asp Thr
 530 535 540
 Gln Met Trp Thr Ile Leu Lys Asp Ser Arg Phe Phe Arg Tyr Leu His
 545 550 555 560
 Thr Ala Val Ile Val Ser Gly Thr Met Leu Val Phe Gly Gly Asn Thr
 565 570 575
 His Asn Asp Thr Ser Met Ser His Gly Ala Lys Cys Phe Ser Ser Asp
 580 585 590
 Phe Met Ala Tyr Asp Ile Ala Cys Asp Arg Trp Ser Val Leu Pro Arg
 595 600 605
 Pro Glu Leu His His Asp Val Asn Arg Phe Gly His Ser Ala Val Leu
 610 615 620
 His Asn Ser Thr Met Tyr Val Phe Gly Gly Phe Asn Ser Leu Leu Leu
 625 630 635 640

Ser Asp Val Leu Val Phe Thr Ser Glu Gln Cys Asp Ala His Arg Ser
 645 650 655
 Glu Ala Ala Cys Val Ala Ala Gly Pro Gly Ile Arg Cys Leu Trp Asp
 660 665 670
 Thr Gln Ser Ser Arg Cys Thr Ser Trp Glu Leu Ala Thr Glu Glu Gln
 675 680 685
 Ala Glu Lys Leu Lys Ser Glu Cys Phe Ser Lys Arg Thr Leu Asp His
 690 695 700
 Asp Arg Cys Asp Gln His Thr Asp Cys Tyr Ser Cys Thr Ala Asn Thr
 705 710 715 720
 Asn Asp Cys His Trp Cys Asn Asp His Cys Val Pro Val Asn His Ser
 725 730 735
 Cys Thr Glu Gly Gln Ile Ser Ile Ala Lys Tyr Asp Asn Cys Pro Lys
 740 745 750
 Asp Asn Pro Met Tyr Tyr Cys Asn Lys Lys Thr Ser Cys Arg Ser Cys
 755 760 765
 Ala Leu Asp Gln Asn Cys Gln Trp Glu Pro Arg Asn Gln Glu Cys Ile
 770 775 780
 Ala Leu Pro Glu Asn Ile Cys Gly Ile Gly Trp His Leu Val Gly Asn
 785 790 795 800
 Ser Cys Leu Lys Ile Thr Thr Ala Lys Glu Asn Tyr Asp Asn Ala Lys
 805 810 815
 Leu Ser Cys Arg Asn His Asn Ala Phe Leu Ala Ser Leu Thr Ser Gln
 820 825 830
 Lys Lys Val Glu Phe Val Leu Lys Gln Leu Arg Leu Met Gln Ser Ser
 835 840 845
 Gln Ser Thr Ser Lys Leu Thr Leu Thr Pro Trp Val Gly Leu Arg Lys
 850 855 860
 Ile Asn Val Ser Tyr Trp Cys Trp Glu Asp Met Ser Pro Phe Thr Asn
 865 870 875 880
 Ser Leu Leu Gln Trp Met Pro Ser Glu Pro Ser Asp Ala Gly Phe Cys
 885 890 895
 Gly Ile Leu Ser Glu Pro Ser Thr Arg Gly Leu Lys Ala Ala Thr Cys
 900 905 910
 Ile Asn Pro Leu Asn Gly Ser Val Cys Glu Arg Pro Ala Asn His Ser
 915 920 925
 Ala Lys Gln Cys Arg Thr Pro Cys Ala Leu Arg Thr Ala Cys Gly Glu
 930 935 940

Cys Thr Ser Ser Ser Ser Glu Cys Met Trp Cys Ser Asn Met Lys Gln
 945 950 955 960
 Cys Val Asp Ser Asn Ala Tyr Val Ala Ser Phe Pro Phe Gly Gln Cys
 965 970 975
 Met Glu Trp Tyr Thr Met Ser Ser Cys Pro Pro Glu Asn Cys Ser Gly
 980 985 990
 Tyr Cys Thr Cys Ser His Cys Leu Glu Gln Pro Gly Cys Gly Trp Cys
 995 1000 1005
 Thr Asp Pro Ser Asn Thr Gly Lys Gly Lys Cys Ile Glu Gly Ser Tyr
 1010 1015 1020
 Lys Gly Pro Val Lys Met Pro Ser His Ala Ser Thr Gly Asn Val Tyr
 1025 1030 1035 1040
 Pro Gln Pro Leu Leu Asn Ser Ser Met Cys Leu Glu Asp Ser Arg Tyr
 1045 1050 1055
 Asn Trp Ser Phe Ile His Cys Pro Ala Cys Gln Cys Asn Gly His Ser
 1060 1065 1070
 Lys Cys Ile Asn Gln Ser Ile Cys Glu Lys Cys Glu Asp Leu Thr Thr
 1075 1080 1085
 Gly Lys His Cys Glu Thr Cys Ile Ser Gly Phe Tyr Gly Asp Pro Thr
 1090 1095 1100
 Asn Gly Gly Lys Cys Gln Pro Cys Lys Cys Asn Gly His Ala Ser Leu
 1105 1110 1115 1120
 Cys Asn Thr Asn Thr Gly Lys Cys Phe Cys Thr Thr Lys Gly Val Lys
 1125 1130 1135
 Gly Glu Glu Cys Gln Leu Cys Glu Val Glu Asn Arg Tyr Gln Gly Asn
 1140 1145 1150
 Pro Leu Lys Gly Thr Cys Tyr Tyr Thr Leu Leu Ile Asp Tyr Gln Phe
 1155 1160 1165
 Thr Phe Ser Leu Ser Gln Glu Asp Asp Arg Tyr Tyr Thr Ala Ile Asn
 1170 1175 1180
 Phe Val Ala Thr Pro Asp Glu Gln Asn Arg Asp Leu Asp Met Phe Ile
 1185 1190 1195 1200
 Asn Ala Ser Lys Asn Phe Asn Leu Asn Ile Thr Trp Ala Thr Ser Phe
 1205 1210 1215
 Pro Ala Gly Thr Gln Thr Gly Glu Glu Val Pro Val Val Ser Lys Thr
 1220 1225 1230
 Asn Ile Lys Glu Tyr Lys Asp Ser Phe Ser Asn Glu Lys Phe Asp Phe
 1235 1240 1245

Arg Asn His Pro Asn Ile Thr Phe Phe Val Tyr Val Ser Asn Phe Thr
 1250 1255 1260

Trp Pro Ile Lys Ile Gln Val Arg Val Thr Ser
 1265 1270 1275

<210> 37
 <211> 883
 <212> PRT
 <213> Homo sapiens

<400> 37
 Met Leu Glu Met Asn Ala Arg Ser Leu Gln Gln Lys Leu Glu Thr Glu
 1 5 10 15
 Arg Glu Leu Lys Gln Arg Leu Leu Glu Glu Gln Ala Lys Leu Gln Gln
 20 25 30
 Gln Met Asp Leu Gln Lys Asn His Ile Phe Arg Leu Thr Gln Gly Leu
 35 40 45
 Gln Glu Ala Leu Asp Arg Ala Asp Leu Leu Lys Thr Glu Arg Ser Asp
 50 55 60
 Leu Glu Tyr Gln Leu Glu Asn Ile Gln Val Leu Tyr Ser His Glu Lys
 65 70 75 80
 Val Lys Met Glu Gly Thr Ile Ser Gln Gln Thr Lys Leu Ile Asp Phe
 85 90 95
 Leu Gln Ala Lys Met Asp Gln Pro Ala Lys Lys Lys Lys Val Pro Leu
 100 105 110
 Gln Tyr Asn Glu Leu Lys Leu Ala Leu Glu Lys Glu Lys Ala Arg Cys
 115 120 125
 Ala Glu Leu Glu Glu Ala Leu Gln Lys Thr Arg Ile Glu Leu Arg Ser
 130 135 140
 Ala Arg Glu Glu Ala Ala His Arg Lys Ala Thr Asp His Pro His Pro
 145 150 155 160
 Ser Thr Pro Ala Thr Ala Arg Gln Gln Ile Ala Met Ser Ala Ile Val
 165 170 175
 Arg Ser Pro Glu His Gln Pro Ser Ala Met Ser Leu Leu Ala Pro Pro
 180 185 190
 Ser Ser Arg Arg Lys Glu Ser Ser Thr Pro Glu Glu Phe Ser Arg Arg
 195 200 205
 Leu Lys Glu Arg Met His His Asn Ile Pro His Arg Phe Asn Val Gly
 210 215 220
 Leu Asn Met Arg Ala Thr Lys Cys Ala Val Cys Leu Asp Thr Val His
 225 230 235 240

Phe Gly Arg Gln Ala Ser Lys Cys Leu Glu Cys Gln Val Met Cys His
245 250 255
Pro Lys Cys Ser Thr Cys Leu Pro Ala Thr Cys Gly Leu Pro Ala Glu
260 265 270
Tyr Ala Thr His Phe Thr Glu Ala Phe Cys Arg Asp Lys Met Asn Ser
275 280 285
Pro Gly Leu Gln Thr Lys Glu Pro Ser Ser Ser Leu His Leu Glu Gly
290 295 300
Trp Met Lys Val Pro Arg Asn Asn Lys Arg Gly Gln Gln Gly Trp Asp
305 310 315 320
Arg Lys Tyr Ile Val Leu Glu Gly Ser Lys Val Leu Ile Tyr Asp Asn
325 330 335
Glu Ala Arg Glu Ala Gly Gln Arg Pro Val Glu Glu Phe Glu Leu Cys
340 345 350
Leu Pro Asp Gly Asp Val Ser Ile His Gly Ala Val Gly Ala Ser Glu
355 360 365
Leu Ala Asn Thr Ala Lys Ala Asp Val Pro Tyr Ile Leu Lys Met Glu
370 375 380
Ser His Pro His Thr Thr Cys Trp Pro Gly Arg Thr Leu Tyr Leu Leu
385 390 395 400
Ala Pro Ser Phe Pro Asp Lys Gln Arg Trp Val Thr Ala Leu Glu Ser
405 410 415
Val Val Ala Gly Gly Arg Val Ser Arg Glu Lys Ala Glu Ala Asp Ala
420 425 430
Lys Leu Leu Gly Asn Ser Leu Leu Lys Leu Glu Gly Asp Asp Arg Leu
435 440 445
Asp Met Asn Cys Thr Leu Pro Phe Ser Asp Gln Val Val Leu Val Gly
450 455 460
Thr Glu Glu Gly Leu Tyr Ala Leu Asn Val Leu Lys Asn Ser Leu Thr
465 470 475 480
His Val Pro Gly Ile Gly Ala Val Phe Gln Ile Tyr Ile Ile Lys Asp
485 490 495
Leu Glu Lys Leu Leu Met Ile Ala Gly Glu Glu Arg Ala Leu Cys Leu
500 505 510
Val Asp Val Lys Lys Val Lys Gln Ser Leu Ala Gln Ser His Leu Pro
515 520 525
Ala Gln Pro Asp Ile Ser Pro Asn Ile Phe Glu Ala Val Lys Gly Cys
530 535 540

His Leu Phe Gly Ala Gly Lys Ile Glu Asn Gly Leu Cys Ile Cys Ala
 545 550 555 560
 Ala Met Pro Ser Lys Val Val Ile Leu Arg Tyr Asn Glu Asn Leu Ser
 565 570 575
 Lys Tyr Cys Ile Arg Lys Glu Ile Glu Thr Ser Glu Pro Cys Ser Cys
 580 585 590
 Ile His Phe Thr Asn Tyr Ser Ile Leu Ile Gly Thr Asn Lys Phe Tyr
 595 600 605
 Glu Ile Asp Met Lys Gln Tyr Thr Leu Glu Glu Phe Leu Asp Lys Asn
 610 615 620
 Asp His Ser Leu Ala Pro Ala Val Phe Ala Ala Ser Ser Asn Ser Phe
 625 630 635 640
 Pro Val Ser Ile Val Gln Val Asn Ser Ala Gly Gln Arg Glu Glu Tyr
 645 650 655
 Leu Leu Cys Phe His Glu Phe Gly Val Phe Val Asp Ser Tyr Gly Arg
 660 665 670
 Arg Ser Arg Thr Asp Asp Leu Lys Trp Ser Arg Leu Pro Leu Ala Phe
 675 680 685
 Ala Tyr Arg Glu Pro Tyr Leu Phe Val Thr His Phe Asn Ser Leu Glu
 690 695 700
 Val Ile Glu Ile Gln Ala Arg Ser Ser Ala Gly Thr Pro Ala Arg Ala
 705 710 715 720
 Tyr Leu Asp Ile Pro Asn Pro Arg Tyr Leu Gly Pro Ala Ile Ser Ser
 725 730 735
 Gly Ala Ile Tyr Leu Ala Ser Ser Tyr Gln Asp Lys Leu Arg Val Ile
 740 745 750
 Cys Cys Lys Gly Asn Leu Val Lys Glu Ser Gly Thr Glu His His Arg
 755 760 765
 Gly Pro Ser Thr Ser Arg Ser Ser Pro Asn Lys Arg Gly Pro Pro Thr
 770 775 780
 Tyr Asn Glu His Ile Thr Lys Arg Val Ala Ser Ser Pro Ala Pro Pro
 785 790 795 800
 Glu Gly Pro Ser His Pro Arg Glu Pro Ser Thr Pro His Arg Tyr Arg
 805 810 815
 Glu Gly Arg Thr Glu Leu Arg Arg Asp Lys Ser Pro Gly Arg Pro Leu
 820 825 830
 Glu Arg Glu Lys Ser Pro Gly Arg Met Leu Ser Thr Arg Arg Glu Arg
 835 840 845

Ser Pro Gly Arg Leu Phe Glu Asp Ser Ser Arg Gly Arg Leu Pro Ala
850 855 860

Gly Ala Val Arg Thr Pro Leu Ser Gln Val Asn Lys Val Trp Asp Gln
865 870 875 880

Ser Ser Val

<210> 38
<211> 1286
<212> PRT
<213> Homo sapiens

<400> 38
Val Leu Asp Asn Gln Ile Lys Lys Asp Leu Ala Asp Lys Glu Thr Leu
1 5 10 15

Glu Asn Met Met Gln Arg His Glu Glu Glu Ala His Glu Lys Gly Lys
20 25 30

Ile Leu Ser Glu Gln Lys Ala Met Ile Asn Ala Met Asp Ser Lys Ile
35 40 45

Arg Ser Leu Glu Gln Arg Ile Val Glu Leu Ser Glu Ala Asn Lys Leu
50 55 60

Ala Ala Asn Ser Ser Leu Phe Thr Gln Arg Asn Met Lys Ala Gln Glu
65 70 75 80

Glu Met Ile Ser Glu Leu Arg Gln Gln Lys Phe Tyr Leu Glu Thr Gln
85 90 95

Ala Gly Lys Leu Glu Ala Gln Asn Arg Lys Leu Glu Glu Gln Leu Glu
100 105 110

Lys Ile Ser His Gln Asp His Ser Asp Lys Asn Arg Leu Leu Glu Leu
115 120 125

Glu Thr Arg Leu Arg Glu Val Ser Leu Glu His Glu Glu Gln Lys Leu
130 135 140

Glu Leu Lys Arg Gln Leu Thr Glu Leu Gln Leu Ser Leu Gln Glu Arg
145 150 155 160

Glu Ser Gln Leu Thr Ala Leu Gln Ala Ala Arg Ala Ala Leu Glu Ser
165 170 175

Gln Leu Arg Gln Ala Lys Thr Glu Leu Glu Glu Thr Thr Ala Glu Ala
180 185 190

Glu Glu Glu Ile Gln Ala Leu Thr Ala His Arg Asp Glu Ile Gln Arg
195 200 205

Lys Phe Asp Ala Leu Arg Asn Ser Cys Thr Val Ile Thr Asp Leu Glu

210					215					220					
Glu	Gln	Leu	Asn	Gln	Leu	Thr	Glu	Asp	Asn	Ala	Glu	Leu	Asn	Asn	Gln
225					230					235					240
Asn	Phe	Tyr	Leu	Ser	Lys	Gln	Leu	Asp	Glu	Ala	Ser	Gly	Ala	Asn	Asp
				245					250					255	
Glu	Ile	Val	Gln	Leu	Arg	Ser	Glu	Val	Asp	His	Leu	Arg	Arg	Glu	Ile
			260					265					270		
Thr	Glu	Arg	Glu	Met	Gln	Leu	Thr	Ser	Gln	Lys	Gln	Thr	Met	Glu	Ala
			275					280					285		
Leu	Lys	Thr	Thr	Cys	Thr	Met	Leu	Glu	Glu	Gln	Val	Met	Asp	Leu	Glu
	290					295					300				
Ala	Leu	Asn	Asp	Glu	Leu	Leu	Glu	Lys	Glu	Arg	Gln	Trp	Glu	Ala	Trp
305					310					315					320
Arg	Ser	Val	Leu	Gly	Asp	Glu	Lys	Ser	Gln	Phe	Glu	Cys	Arg	Val	Arg
				325					330					335	
Glu	Leu	Gln	Arg	Met	Leu	Asp	Thr	Glu	Lys	Gln	Ser	Arg	Ala	Arg	Ala
			340					345					350		
Asp	Gln	Arg	Ile	Thr	Glu	Ser	Arg	Gln	Val	Val	Glu	Leu	Ala	Val	Lys
			355					360					365		
Glu	His	Lys	Ala	Glu	Ile	Leu	Ala	Leu	Gln	Gln	Ala	Leu	Lys	Glu	Gln
	370					375					380				
Lys	Leu	Lys	Ala	Glu	Ser	Leu	Ser	Asp	Lys	Leu	Asn	Asp	Leu	Glu	Lys
385						390					395				400
Lys	His	Ala	Met	Leu	Glu	Met	Asn	Ala	Arg	Ser	Leu	Gln	Gln	Lys	Leu
			405						410					415	
Glu	Thr	Glu	Arg	Glu	Leu	Lys	Gln	Arg	Leu	Leu	Glu	Glu	Gln	Ala	Lys
			420					425					430		
Leu	Gln	Gln	Gln	Met	Asp	Leu	Gln	Lys	Asn	His	Ile	Phe	Arg	Leu	Thr
			435					440					445		
Gln	Gly	Leu	Gln	Glu	Ala	Leu	Asp	Arg	Ala	Asp	Leu	Leu	Lys	Thr	Glu
	450					455					460				
Arg	Ser	Asp	Leu	Glu	Tyr	Gln	Leu	Glu	Asn	Ile	Gln	Val	Leu	Tyr	Ser
465					470					475					480
His	Glu	Lys	Val	Lys	Met	Glu	Gly	Thr	Ile	Ser	Gln	Gln	Thr	Lys	Leu
				485					490					495	
Ile	Asp	Phe	Leu	Gln	Ala	Lys	Met	Asp	Gln	Pro	Ala	Lys	Lys	Lys	Lys
			500					505					510		
Val	Pro	Leu	Gln	Tyr	Asn	Glu	Leu	Lys	Leu	Ala	Leu	Glu	Lys	Glu	Lys

515					520					525						
Ala	Arg	Cys	Ala	Glu	Leu	Glu	Glu	Ala	Leu	Gln	Lys	Thr	Arg	Ile	Glu	
530					535					540						
Leu	Arg	Ser	Ala	Arg	Glu	Glu	Ala	Ala	His	Arg	Lys	Ala	Thr	Asp	His	
545					550					555					560	
Pro	His	Pro	Ser	Thr	Pro	Ala	Thr	Ala	Arg	Gln	Gln	Ile	Ala	Met	Ser	
565					570					575						
Ala	Ile	Val	Arg	Ser	Pro	Glu	His	Gln	Pro	Ser	Ala	Met	Ser	Leu	Leu	
580					585					590						
Ala	Pro	Pro	Ser	Ser	Arg	Arg	Lys	Glu	Ser	Ser	Thr	Pro	Glu	Glu	Phe	
595					600					605						
Ser	Arg	Arg	Leu	Lys	Glu	Arg	Met	His	His	Asn	Ile	Pro	His	Arg	Phe	
610					615					620						
Asn	Val	Gly	Leu	Asn	Met	Arg	Ala	Thr	Lys	Cys	Ala	Val	Cys	Leu	Asp	
625					630					635					640	
Thr	Val	His	Phe	Gly	Arg	Gln	Ala	Ser	Lys	Cys	Leu	Glu	Cys	Gln	Val	
645					650					655						
Met	Cys	His	Pro	Lys	Cys	Ser	Thr	Cys	Leu	Pro	Ala	Thr	Cys	Gly	Leu	
660					665					670						
Pro	Ala	Glu	Tyr	Ala	Thr	His	Phe	Thr	Glu	Ala	Phe	Cys	Arg	Asp	Lys	
675					680					685						
Met	Asn	Ser	Pro	Gly	Leu	Gln	Thr	Lys	Glu	Pro	Ser	Ser	Ser	Leu	His	
690					695					700						
Leu	Glu	Gly	Trp	Met	Lys	Val	Pro	Arg	Asn	Asn	Lys	Arg	Gly	Gln	Gln	
705					710					715					720	
Gly	Trp	Asp	Arg	Lys	Tyr	Ile	Val	Leu	Glu	Gly	Ser	Lys	Val	Leu	Ile	
725					730					735						
Tyr	Asp	Asn	Glu	Ala	Arg	Glu	Ala	Gly	Gln	Arg	Pro	Val	Glu	Glu	Phe	
740					745					750						
Glu	Leu	Cys	Leu	Pro	Asp	Gly	Asp	Val	Ser	Ile	His	Gly	Ala	Val	Gly	
755					760					765						
Ala	Ser	Glu	Leu	Ala	Asn	Thr	Ala	Lys	Ala	Asp	Val	Pro	Tyr	Ile	Leu	
770					775					780						
Lys	Met	Glu	Ser	His	Pro	His	Thr	Thr	Cys	Trp	Pro	Gly	Arg	Thr	Leu	
785					790					795					800	
Tyr	Leu	Leu	Ala	Pro	Ser	Phe	Pro	Asp	Lys	Gln	Arg	Trp	Val	Thr	Ala	
805					810					815						
Leu	Glu	Ser	Val	Val	Ala	Gly	Gly	Arg	Val	Ser	Arg	Glu	Lys	Ala	Glu	

820					825					830						
Ala	Asp	Ala	Lys	Leu	Leu	Gly	Asn	Ser	Leu	Leu	Lys	Leu	Glu	Gly	Asp	
835					840					845						
Asp	Arg	Leu	Asp	Met	Asn	Cys	Thr	Leu	Pro	Phe	Ser	Asp	Gln	Val	Val	
850					855					860						
Leu	Val	Gly	Thr	Glu	Glu	Gly	Leu	Tyr	Ala	Leu	Asn	Val	Leu	Lys	Asn	
865					870					875					880	
Ser	Leu	Thr	His	Val	Pro	Gly	Ile	Gly	Ala	Val	Phe	Gln	Ile	Tyr	Ile	
885					890					895						
Ile	Lys	Asp	Leu	Glu	Lys	Leu	Leu	Met	Ile	Ala	Gly	Glu	Glu	Arg	Ala	
900					905					910						
Leu	Cys	Leu	Val	Asp	Val	Lys	Lys	Val	Lys	Gln	Ser	Leu	Ala	Gln	Ser	
915					920					925						
His	Leu	Pro	Ala	Gln	Pro	Asp	Ile	Ser	Pro	Asn	Ile	Phe	Glu	Ala	Val	
930					935					940						
Lys	Gly	Cys	His	Leu	Phe	Gly	Ala	Gly	Lys	Ile	Glu	Asn	Gly	Leu	Cys	
945					950					955					960	
Ile	Cys	Ala	Ala	Met	Pro	Ser	Lys	Val	Val	Ile	Leu	Arg	Tyr	Asn	Glu	
965					970					975						
Asn	Leu	Ser	Lys	Tyr	Cys	Ile	Arg	Lys	Glu	Ile	Glu	Thr	Ser	Glu	Pro	
980					985					990						
Cys	Ser	Cys	Ile	His	Phe	Thr	Asn	Tyr	Ser	Ile	Leu	Ile	Gly	Thr	Asn	
995					1000					1005						
Lys	Phe	Tyr	Glu	Ile	Asp	Met	Lys	Gln	Tyr	Thr	Leu	Glu	Glu	Phe	Leu	
1010					1015					1020						
Asp	Lys	Asn	Asp	His	Ser	Leu	Ala	Pro	Ala	Val	Phe	Ala	Ala	Ser	Ser	
1025					1030					1035					1040	
Asn	Ser	Phe	Pro	Val	Ser	Ile	Val	Gln	Val	Asn	Ser	Ala	Gly	Gln	Arg	
1045					1050					1055						
Glu	Glu	Tyr	Leu	Leu	Cys	Phe	His	Glu	Phe	Gly	Val	Phe	Val	Asp	Ser	
1060					1065					1070						
Tyr	Gly	Arg	Arg	Ser	Arg	Thr	Asp	Asp	Leu	Lys	Trp	Ser	Arg	Leu	Pro	
1075					1080					1085						
Leu	Ala	Phe	Ala	Tyr	Arg	Glu	Pro	Tyr	Leu	Phe	Val	Thr	His	Phe	Asn	
1090					1095					1100						
Ser	Leu	Glu	Val	Ile	Glu	Ile	Gln	Ala	Arg	Ser	Ser	Ala	Gly	Thr	Pro	
1105					1110					1115					1120	
Ala	Arg	Ala	Tyr	Leu	Asp	Ile	Pro	Asn	Pro	Arg	Tyr	Leu	Gly	Pro	Ala	

1125	1130	1135
Ile Ser Ser Gly Ala Ile Tyr Leu Ala Ser Ser Tyr Gln Asp Lys Leu		
1140	1145	1150
Arg Val Ile Cys Cys Lys Gly Asn Leu Val Lys Glu Ser Gly Thr Glu		
1155	1160	1165
His His Arg Gly Pro Ser Thr Ser Arg Ser Ser Pro Asn Lys Arg Gly		
1170	1175	1180
Pro Pro Thr Tyr Asn Glu His Ile Thr Lys Arg Val Ala Ser Ser Pro		
1185	1190	1200
Ala Pro Pro Glu Gly Pro Ser His Pro Arg Glu Pro Ser Thr Pro His		
1205	1210	1215
Arg Tyr Arg Glu Gly Arg Thr Glu Leu Arg Arg Asp Lys Ser Pro Gly		
1220	1225	1230
Arg Pro Leu Glu Arg Glu Lys Ser Pro Gly Arg Met Leu Ser Thr Arg		
1235	1240	1245
Arg Glu Arg Ser Pro Gly Arg Leu Phe Glu Asp Ser Ser Arg Gly Arg		
1250	1255	1260
Leu Pro Ala Gly Ala Val Arg Thr Pro Leu Ser Gln Val Asn Lys Val		
1265	1270	1280
Trp Asp Gln Ser Ser Val		
1285		

<210> 39
 <211> 940
 <212> PRT
 <213> Homo sapiens

<400> 39
Gln Ser Arg Ala Arg Ala Asp Gln Arg Ile Thr Glu Ser Arg Gln Val
1 5 10 15
Val Glu Leu Ala Val Lys Glu His Lys Ala Glu Ile Leu Ala Leu Gln
20 25 30
Gln Ala Leu Lys Glu Gln Lys Leu Lys Ala Glu Ser Leu Ser Asp Lys
35 40 45
Leu Asn Asp Leu Glu Lys Lys His Ala Met Leu Glu Met Asn Ala Arg
50 55 60
Ser Leu Gln Gln Lys Leu Glu Thr Glu Arg Glu Leu Lys Gln Arg Leu
65 70 75 80
Leu Glu Glu Gln Ala Lys Leu Gln Gln Gln Met Asp Leu Gln Lys Asn
85 90 95

His Ile Phe Arg Leu Thr Gln Gly Leu Gln Glu Ala Leu Asp Arg Ala
 100 105 110
 Asp Leu Leu Lys Thr Glu Arg Ser Asp Leu Glu Tyr Gln Leu Glu Asn
 115 120 125
 Ile Gln Val Leu Tyr Ser His Glu Lys Val Lys Met Glu Gly Thr Ile
 130 135 140
 Ser Gln Gln Thr Lys Leu Ile Asp Phe Leu Gln Ala Lys Met Asp Gln
 145 150 155 160
 Pro Ala Lys Lys Lys Lys Val Pro Leu Gln Tyr Asn Glu Leu Lys Leu
 165 170 175
 Ala Leu Glu Lys Glu Lys Ala Arg Cys Ala Glu Leu Glu Glu Ala Leu
 180 185 190
 Gln Lys Thr Arg Ile Glu Leu Arg Ser Ala Arg Glu Glu Ala Ala His
 195 200 205
 Arg Lys Ala Thr Asp His Pro His Pro Ser Thr Pro Ala Thr Ala Arg
 210 215 220
 Gln Gln Ile Ala Met Ser Ala Ile Val Arg Ser Pro Glu His Gln Pro
 225 230 235 240
 Ser Ala Met Ser Leu Leu Ala Pro Pro Ser Ser Arg Arg Lys Glu Ser
 245 250 255
 Ser Thr Pro Glu Glu Phe Ser Arg Arg Leu Lys Glu Arg Met His His
 260 265 270
 Asn Ile Pro His Arg Phe Asn Val Gly Leu Asn Met Arg Ala Thr Lys
 275 280 285
 Cys Ala Val Cys Leu Asp Thr Val His Phe Gly Arg Gln Ala Ser Lys
 290 295 300
 Cys Leu Glu Cys Gln Val Met Cys His Pro Lys Cys Ser Thr Cys Leu
 305 310 315 320
 Pro Ala Thr Cys Gly Leu Pro Ala Glu Tyr Ala Thr His Phe Thr Glu
 325 330 335
 Ala Phe Cys Arg Asp Lys Met Asn Ser Pro Gly Leu Gln Thr Lys Glu
 340 345 350
 Pro Ser Ser Ser Leu His Leu Glu Gly Trp Met Lys Val Pro Arg Asn
 355 360 365
 Asn Lys Arg Gly Gln Gln Gly Trp Asp Arg Lys Tyr Ile Val Leu Glu
 370 375 380
 Gly Ser Lys Val Leu Ile Tyr Asp Asn Glu Ala Arg Glu Ala Gly Gln
 385 390 395 400

Arg Pro Val Glu Glu Phe Glu Leu Cys Leu Pro Asp Gly Asp Val Ser
 405 410 415
 Ile His Gly Ala Val Gly Ala Ser Glu Leu Ala Asn Thr Ala Lys Ala
 420 425 430
 Asp Val Pro Tyr Ile Leu Lys Met Glu Ser His Pro His Thr Thr Cys
 435 440 445
 Trp Pro Gly Arg Thr Leu Tyr Leu Leu Ala Pro Ser Phe Pro Asp Lys
 450 455 460
 Gln Arg Trp Val Thr Ala Leu Glu Ser Val Val Ala Gly Gly Arg Val
 465 470 475 480
 Ser Arg Glu Lys Ala Glu Ala Asp Ala Lys Leu Leu Gly Asn Ser Leu
 485 490 495
 Leu Lys Leu Glu Gly Asp Asp Arg Leu Asp Met Asn Cys Thr Leu Pro
 500 505 510
 Phe Ser Asp Gln Val Val Leu Val Gly Thr Glu Glu Gly Leu Tyr Ala
 515 520 525
 Leu Asn Val Leu Lys Asn Ser Leu Thr His Val Pro Gly Ile Gly Ala
 530 535 540
 Val Phe Gln Ile Tyr Ile Ile Lys Asp Leu Glu Lys Leu Leu Met Ile
 545 550 555 560
 Ala Gly Glu Glu Arg Ala Leu Cys Leu Val Asp Val Lys Lys Val Lys
 565 570 575
 Gln Ser Leu Ala Gln Ser His Leu Pro Ala Gln Pro Asp Ile Ser Pro
 580 585 590
 Asn Ile Phe Glu Ala Val Lys Gly Cys His Leu Phe Gly Ala Gly Lys
 595 600 605
 Ile Glu Asn Gly Leu Cys Ile Cys Ala Ala Met Pro Ser Lys Val Val
 610 615 620
 Ile Leu Arg Tyr Asn Glu Asn Leu Ser Lys Tyr Cys Ile Arg Lys Glu
 625 630 635 640
 Ile Glu Thr Ser Glu Pro Cys Ser Cys Ile His Phe Thr Asn Tyr Ser
 645 650 655
 Ile Leu Ile Gly Thr Asn Lys Phe Tyr Glu Ile Asp Met Lys Gln Tyr
 660 665 670
 Thr Leu Glu Glu Phe Leu Asp Lys Asn Asp His Ser Leu Ala Pro Ala
 675 680 685
 Val Phe Ala Ala Ser Ser Asn Ser Phe Pro Val Ser Ile Val Gln Val
 690 695 700

Asn Ser Ala Gly Gln Arg Glu Glu Tyr Leu Leu Cys Phe His Glu Phe
 705 710 715 720
 Gly Val Phe Val Asp Ser Tyr Gly Arg Arg Ser Arg Thr Asp Asp Leu
 725 730 735
 Lys Trp Ser Arg Leu Pro Leu Ala Phe Ala Tyr Arg Glu Pro Tyr Leu
 740 745 750
 Phe Val Thr His Phe Asn Ser Leu Glu Val Ile Glu Ile Gln Ala Arg
 755 760 765
 Ser Ser Ala Gly Thr Pro Ala Arg Ala Tyr Leu Asp Ile Pro Asn Pro
 770 775 780
 Arg Tyr Leu Gly Pro Ala Ile Ser Ser Gly Ala Ile Tyr Leu Ala Ser
 785 790 795 800
 Ser Tyr Gln Asp Lys Leu Arg Val Ile Cys Cys Lys Gly Asn Leu Val
 805 810 815
 Lys Glu Ser Gly Thr Glu His His Arg Gly Pro Ser Thr Ser Arg Ser
 820 825 830
 Ser Pro Asn Lys Arg Gly Pro Pro Thr Tyr Asn Glu His Ile Thr Lys
 835 840 845
 Arg Val Ala Ser Ser Pro Ala Pro Pro Glu Gly Pro Ser His Pro Arg
 850 855 860
 Glu Pro Ser Thr Pro His Arg Tyr Arg Glu Gly Arg Thr Glu Leu Arg
 865 870 875 880
 Arg Asp Lys Ser Pro Gly Arg Pro Leu Glu Arg Glu Lys Ser Pro Gly
 885 890 895
 Arg Met Leu Ser Thr Arg Arg Glu Arg Ser Pro Gly Arg Leu Phe Glu
 900 905 910
 Asp Ser Ser Arg Gly Arg Leu Pro Ala Gly Ala Val Arg Thr Pro Leu
 915 920 925
 Ser Gln Val Asn Lys Val Trp Asp Gln Ser Ser Val
 930 935 940

<210> 40

<211> 1641

<212> PRT

<213> Mus musculus

<400> 40

Pro Phe Val Pro Thr Leu Lys Ser Asp Asp Asp Thr Ser Asn Phe Asp
 1 5 10 15

Glu Pro Glu Lys Asn Ser Trp Val Ser Ser Ser Val Cys Gln Leu Ser
 20 25 30

Pro Ser Gly Phe Ser Gly Glu Glu Leu Pro Phe Val Gly Phe Ser Tyr
 35 40 45
 Ser Lys Ala Leu Gly Tyr Leu Gly Arg Ser Glu Ser Val Val Ser Ser
 50 55 60
 Leu Asp Ser Pro Ala Lys Val Ser Ser Met Glu Lys Lys Leu Leu Ile
 65 70 75 80
 Lys Ser Lys Glu Leu Gln Asp Ser Gln Asp Lys Cys His Lys Met Glu
 85 90 95
 Gln Glu Met Thr Arg Leu His Arg Arg Val Ser Glu Val Glu Ala Val
 100 105 110
 Leu Ser Gln Lys Glu Val Glu Leu Lys Ala Ser Glu Thr Gln Arg Ser
 115 120 125
 Leu Leu Glu Gln Asp Leu Ala Thr Tyr Ile Thr Glu Cys Ser Ser Leu
 130 135 140
 Lys Arg Ser Leu Glu Gln Ala Arg Met Glu Val Ser Gln Glu Asp Asp
 145 150 155 160
 Lys Ala Leu Gln Leu Leu His Asp Ile Arg Glu Gln Ser Arg Lys Leu
 165 170 175
 Gln Glu Ile Lys Glu Gln Glu Tyr Gln Ala Gln Val Glu Glu Met Arg
 180 185 190
 Leu Met Met Asn Gln Leu Glu Glu Asp Leu Val Ser Ala Arg Arg Arg
 195 200 205
 Ser Asp Leu Tyr Glu Ser Glu Leu Arg Glu Ser Arg Leu Ala Ala Glu
 210 215 220
 Glu Phe Lys Arg Lys Ala Asn Glu Cys Gln His Lys Leu Met Lys Ala
 225 230 235 240
 Lys Asp Gln Gly Lys Pro Glu Val Gly Glu Tyr Ser Lys Leu Glu Lys
 245 250 255
 Ile Asn Ala Glu Gln Gln Leu Lys Ile Gln Glu Leu Gln Glu Lys Leu
 260 265 270
 Glu Lys Ala Val Lys Ala Ser Thr Glu Ala Thr Glu Leu Leu Gln Asn
 275 280 285
 Ile Arg Gln Ala Lys Glu Arg Ala Glu Arg Glu Leu Glu Lys Leu His
 290 295 300
 Asn Arg Glu Asp Ser Ser Glu Gly Ile Lys Lys Lys Leu Val Glu Ala
 305 310 315 320
 Glu Glu Leu Glu Glu Lys His Arg Glu Ala Gln Val Ser Ala Gln His
 325 330 335

Leu Glu Val His Leu Lys Gln Lys Glu Gln His Tyr Glu Glu Lys Ile
 340 345 350
 Lys Val Leu Asp Asn Gln Ile Lys Lys Asp Leu Ala Asp Lys Glu Ser
 355 360 365
 Leu Glu Asn Met Met Gln Arg His Glu Glu Glu Ala His Glu Lys Gly
 370 375 380
 Lys Ile Leu Ser Glu Gln Lys Ala Met Ile Asn Ala Met Asp Ser Lys
 385 390 395 400
 Ile Arg Ser Leu Glu Gln Arg Ile Val Glu Leu Ser Glu Ala Asn Lys
 405 410 415
 Leu Ala Ala Asn Ser Ser Leu Phe Thr Gln Arg Asn Met Lys Ala Gln
 420 425 430
 Glu Glu Met Ile Ser Glu Leu Arg Gln Gln Lys Phe Tyr Leu Glu Thr
 435 440 445
 Gln Ala Gly Lys Leu Glu Ala Gln Asn Arg Lys Leu Glu Glu Gln Leu
 450 455 460
 Glu Lys Ile Ser His Gln Asp His Ser Asp Lys Ser Arg Leu Leu Glu
 465 470 475 480
 Leu Glu Thr Arg Leu Arg Glu Val Ser Leu Glu His Glu Glu Gln Lys
 485 490 495
 Leu Glu Leu Lys Arg Gln Leu Thr Glu Leu Gln Leu Ser Leu Gln Glu
 500 505 510
 Arg Glu Ser Gln Leu Thr Ala Leu Gln Ala Ala Arg Ala Ala Leu Glu
 515 520 525
 Ser Gln Leu Arg Gln Ala Lys Thr Glu Leu Glu Glu Thr Thr Ala Glu
 530 535 540
 Ala Glu Glu Glu Ile Gln Ala Leu Thr Ala His Arg Asp Glu Ile Gln
 545 550 555 560
 Arg Lys Phe Asp Ala Leu Arg Asn Ser Cys Thr Val Ile Thr Asp Leu
 565 570 575
 Glu Glu Gln Leu Asn Gln Leu Thr Glu Asp Asn Ala Glu Leu Asn Asn
 580 585 590
 Gln Asn Phe Tyr Leu Ser Lys Gln Leu Asp Glu Ala Ser Gly Ala Asn
 595 600 605
 Asp Glu Ile Val Gln Leu Arg Ser Glu Val Asp His Leu Arg Arg Glu
 610 615 620
 Ile Thr Glu Arg Glu Met Gln Leu Thr Ser Gln Lys Gln Thr Met Glu
 625 630 635 640

Ala Leu Lys Thr Thr Cys Thr Met Leu Glu Glu Gln Val Leu Asp Leu
 645 650 655
 Glu Ala Leu Asn Asp Glu Leu Leu Glu Lys Glu Arg Gln Trp Glu Ala
 660 665 670
 Trp Arg Ser Val Leu Gly Asp Glu Lys Ser Gln Phe Glu Cys Arg Val
 675 680 685
 Arg Glu Leu Gln Arg Met Leu Asp Thr Glu Lys Gln Ser Arg Ala Arg
 690 695 700
 Ala Asp Gln Arg Ile Thr Glu Ser Arg Gln Val Val Glu Leu Ala Val
 705 710 715 720
 Lys Glu His Lys Ala Glu Ile Leu Ala Leu Gln Gln Ala Leu Lys Glu
 725 730 735
 Gln Lys Leu Lys Ala Glu Ser Leu Ser Asp Lys Leu Asn Asp Leu Glu
 740 745 750
 Lys Lys His Ala Met Leu Glu Met Asn Ala Arg Ser Leu Gln Gln Lys
 755 760 765
 Leu Glu Thr Glu Arg Glu Leu Lys Gln Arg Leu Leu Glu Glu Gln Ala
 770 775 780
 Lys Leu Gln Gln Gln Met Asp Leu Gln Lys Asn His Ile Phe Arg Leu
 785 790 795 800
 Thr Gln Gly Leu Gln Glu Ala Leu Asp Arg Ala Asp Leu Leu Lys Thr
 805 810 815
 Glu Arg Ser Asp Leu Glu Tyr Gln Leu Glu Asn Ile Gln Val Leu Tyr
 820 825 830
 Ser His Glu Lys Val Lys Met Glu Gly Thr Ile Ser Gln Gln Thr Lys
 835 840 845
 Leu Ile Asp Phe Leu Gln Ala Lys Met Asp Gln Pro Ala Lys Lys Lys
 850 855 860
 Lys Val Pro Leu Gln Tyr Asn Glu Leu Lys Leu Ala Leu Glu Lys Glu
 865 870 875 880
 Lys Ala Arg Cys Ala Glu Leu Glu Glu Ala Leu Gln Lys Thr Arg Ile
 885 890 895
 Glu Leu Arg Ser Ala Arg Glu Glu Ala Ala His Arg Lys Ala Thr Asp
 900 905 910
 His Pro His Pro Ser Thr Pro Ala Thr Ala Arg Gln Gln Ile Ala Met
 915 920 925
 Ser Ala Ile Val Arg Ser Pro Glu His Gln Pro Ser Ala Met Ser Leu
 930 935 940

Leu Ala Pro Pro Ser Ser Arg Arg Lys Glu Ser Ser Thr Pro Glu Glu
 945 950 955 960
 Phe Ser Arg Arg Leu Lys Glu Arg Met His His Asn Ile Pro His Arg
 965 970 975
 Phe Asn Val Gly Leu Asn Met Arg Ala Thr Lys Cys Ala Val Cys Leu
 980 985 990
 Asp Thr Val His Phe Gly Arg Gln Ala Ser Lys Cys Leu Glu Cys Gln
 995 1000 1005
 Val Met Cys His Pro Lys Cys Ser Thr Cys Leu Pro Ala Thr Cys Gly
 1010 1015 1020
 Leu Pro Ala Glu Tyr Ala Thr His Phe Thr Glu Ala Phe Cys Arg Asp
 1025 1030 1035 1040
 Lys Met Asn Ser Pro Gly Leu Gln Ser Lys Glu Pro Gly Ser Ser Leu
 1045 1050 1055
 His Leu Glu Gly Trp Met Lys Val Pro Arg Asn Asn Lys Arg Gly Gln
 1060 1065 1070
 Gln Gly Trp Asp Arg Lys Tyr Ile Val Leu Glu Gly Ser Lys Val Leu
 1075 1080 1085
 Ile Tyr Asp Asn Glu Ala Arg Glu Ala Gly Gln Arg Pro Val Glu Glu
 1090 1095 1100
 Phe Glu Leu Cys Leu Pro Asp Gly Asp Val Ser Ile His Gly Ala Val
 1105 1110 1115 1120
 Gly Ala Ser Glu Leu Ala Asn Thr Ala Lys Ala Asp Val Pro Tyr Ile
 1125 1130 1135
 Leu Lys Met Glu Ser His Pro His Thr Thr Cys Trp Pro Gly Arg Thr
 1140 1145 1150
 Leu Tyr Leu Leu Ala Pro Ser Phe Pro Asp Lys Gln Arg Trp Val Thr
 1155 1160 1165
 Ala Leu Glu Ser Val Val Ala Gly Gly Arg Val Ser Arg Glu Lys Ala
 1170 1175 1180
 Glu Ala Asp Ala Lys Leu Leu Gly Asn Ser Leu Leu Lys Leu Glu Gly
 1185 1190 1195 1200
 Asp Asp Arg Leu Asp Met Asn Cys Thr Leu Pro Phe Ser Asp Gln Val
 1205 1210 1215
 Val Leu Val Gly Thr Glu Glu Gly Leu Tyr Ala Leu Asn Val Leu Lys
 1220 1225 1230
 Asn Ser Leu Thr His Ile Pro Gly Ile Gly Ala Val Phe Gln Ile Tyr
 1235 1240 1245

Ile Ile Lys Asp Leu Glu Lys Leu Leu Met Ile Ala Gly Glu Glu Arg
 1250 1255 1260
 Ala Leu Cys Leu Val Asp Val Lys Lys Val Lys Gln Ser Leu Ala Gln
 1265 1270 1275 1280
 Ser His Leu Pro Ala Gln Pro Asp Val Ser Pro Asn Ile Phe Glu Ala
 1285 1290 1295
 Val Lys Gly Cys His Leu Phe Ala Ala Gly Lys Ile Glu Asn Ser Leu
 1300 1305 1310
 Cys Ile Cys Ala Ala Met Pro Ser Lys Val Val Ile Leu Arg Tyr Asn
 1315 1320 1325
 Asp Asn Leu Ser Lys Tyr Cys Ile Arg Lys Glu Ile Glu Thr Ser Glu
 1330 1335 1340
 Pro Cys Ser Cys Ile His Phe Thr Asn Tyr Ser Ile Leu Ile Gly Thr
 1345 1350 1355 1360
 Asn Lys Phe Tyr Glu Ile Asp Met Lys Gln Tyr Thr Leu Asp Glu Phe
 1365 1370 1375
 Leu Asp Lys Asn Asp His Ser Leu Ala Pro Ala Val Phe Ala Ser Ser
 1380 1385 1390
 Ser Asn Ser Phe Pro Val Ser Ile Val Gln Ala Asn Ser Ala Gly Gln
 1395 1400 1405
 Arg Glu Glu Tyr Leu Leu Cys Phe His Glu Phe Gly Val Phe Val Asp
 1410 1415 1420
 Ser Tyr Gly Arg Arg Ser Arg Thr Asp Asp Leu Lys Trp Ser Arg Leu
 1425 1430 1435 1440
 Pro Leu Ala Phe Ala Tyr Arg Glu Pro Tyr Leu Phe Val Thr His Phe
 1445 1450 1455
 Asn Ser Leu Glu Val Ile Glu Ile Gln Ala Arg Ser Ser Leu Gly Ser
 1460 1465 1470
 Pro Ala Arg Ala Tyr Leu Glu Ile Pro Asn Pro Arg Tyr Leu Gly Pro
 1475 1480 1485
 Ala Ile Ser Ser Gly Ala Ile Tyr Leu Ala Ser Ser Tyr Gln Asp Lys
 1490 1495 1500
 Leu Arg Val Ile Cys Cys Lys Gly Asn Leu Val Lys Glu Ser Gly Thr
 1505 1510 1515 1520
 Glu Gln His Arg Val Pro Ser Thr Ser Arg Ser Ser Pro Asn Lys Arg
 1525 1530 1535
 Gly Pro Pro Thr Tyr Asn Glu His Ile Thr Lys Arg Val Ala Ser Ser
 1540 1545 1550

Pro Ala Pro Pro Glu Gly Pro Ser His Pro Arg Glu Pro Ser Thr Pro
1555 1560 1565

His Arg Tyr Arg Asp Arg Glu Gly Arg Thr Glu Leu Arg Arg Asp Lys
1570 1575 1580

Ser Pro Gly Arg Pro Leu Glu Arg Glu Lys Ser Pro Gly Arg Met Leu
1585 1590 1595 1600

Ser Thr Arg Arg Glu Arg Ser Pro Gly Arg Leu Phe Glu Asp Ser Ser
1605 1610 1615

Arg Gly Arg Leu Pro Ala Gly Ala Val Arg Thr Pro Leu Ser Gln Val
1620 1625 1630

Asn Lys Val Trp Asp Gln Ser Ser Val
1635 1640

<210> 41
<211> 1597
<212> PRT
<213> Mus musculus

<400> 41
Met Leu Leu Gly Glu Glu Ala Met Met Glu Gln Glu Met Thr Arg Leu
1 5 10 15

His Arg Arg Val Ser Glu Val Glu Ala Val Leu Ser Gln Lys Glu Val
20 25 30

Glu Leu Lys Ala Ser Glu Thr Gln Arg Ser Leu Leu Glu Gln Asp Leu
35 40 45

Ala Thr Tyr Ile Thr Glu Cys Ser Ser Leu Lys Arg Ser Leu Glu Gln
50 55 60

Ala Arg Met Glu Val Ser Gln Glu Asp Asp Lys Ala Leu Gln Leu Leu
65 70 75 80

His Asp Ile Arg Glu Gln Ser Arg Lys Leu Gln Glu Ile Lys Glu Gln
85 90 95

Glu Tyr Gln Ala Gln Val Glu Glu Met Arg Leu Met Met Asn Gln Leu
100 105 110

Glu Glu Asp Leu Val Ser Ala Arg Arg Arg Ser Asp Leu Tyr Glu Ser
115 120 125

Glu Leu Arg Glu Ser Arg Leu Ala Ala Glu Glu Phe Lys Arg Lys Ala
130 135 140

Asn Glu Cys Gln His Lys Leu Met Lys Ala Lys Asp Gln Gly Lys Pro
145 150 155 160

Glu Val Gly Glu Tyr Ser Lys Leu Glu Lys Ile Asn Ala Glu Gln Gln

165					170					175					
Leu	Lys	Ile	Gln	Glu	Leu	Gln	Glu	Lys	Leu	Glu	Lys	Ala	Val	Lys	Ala
			180					185					190		
Ser	Thr	Glu	Ala	Thr	Glu	Leu	Leu	Gln	Asn	Ile	Arg	Gln	Ala	Lys	Glu
		195					200					205			
Arg	Ala	Glu	Arg	Glu	Leu	Glu	Lys	Leu	His	Asn	Arg	Glu	Asp	Ser	Ser
	210					215					220				
Glu	Gly	Ile	Lys	Lys	Lys	Leu	Val	Glu	Ala	Glu	Glu	Arg	Arg	His	Ser
225						230					235				240
Leu	Glu	Asn	Lys	Val	Lys	Arg	Leu	Glu	Thr	Met	Glu	Arg	Arg	Glu	Asn
			245						250					255	
Arg	Leu	Lys	Asp	Asp	Ile	Gln	Thr	Lys	Ser	Glu	Gln	Ile	Gln	Gln	Met
			260					265					270		
Ala	Asp	Lys	Ile	Leu	Glu	Leu	Glu	Glu	Lys	His	Arg	Glu	Ala	Gln	Val
	275						280					285			
Ser	Ala	Gln	His	Leu	Glu	Val	His	Leu	Lys	Gln	Lys	Glu	Gln	His	Tyr
	290					295					300				
Glu	Glu	Lys	Ile	Lys	Val	Leu	Asp	Asn	Gln	Ile	Lys	Lys	Asp	Leu	Ala
305						310					315				320
Asp	Lys	Glu	Ser	Leu	Glu	Asn	Met	Met	Gln	Arg	His	Glu	Glu	Glu	Ala
				325					330					335	
His	Glu	Lys	Gly	Lys	Ile	Leu	Ser	Glu	Gln	Lys	Ala	Met	Ile	Asn	Ala
			340					345					350		
Met	Asp	Ser	Lys	Ile	Arg	Ser	Leu	Glu	Gln	Arg	Ile	Val	Glu	Leu	Ser
	355						360					365			
Glu	Ala	Asn	Lys	Leu	Ala	Ala	Asn	Ser	Ser	Leu	Phe	Thr	Gln	Arg	Asn
	370					375					380				
Met	Lys	Ala	Gln	Glu	Glu	Met	Ile	Ser	Glu	Leu	Arg	Gln	Gln	Lys	Phe
385						390					395				400
Tyr	Leu	Glu	Thr	Gln	Ala	Gly	Lys	Leu	Glu	Ala	Gln	Asn	Arg	Lys	Leu
			405						410					415	
Glu	Glu	Gln	Leu	Glu	Lys	Ile	Ser	His	Gln	Asp	His	Ser	Asp	Lys	Ser
			420					425					430		
Arg	Leu	Leu	Glu	Leu	Glu	Thr	Arg	Leu	Arg	Glu	Val	Ser	Leu	Glu	His
	435						440					445			
Glu	Glu	Gln	Lys	Leu	Glu	Leu	Lys	Arg	Gln	Leu	Thr	Glu	Leu	Gln	Leu
	450					455					460				
Ser	Leu	Gln	Glu	Arg	Glu	Ser	Gln	Leu	Thr	Ala	Leu	Gln	Ala	Ala	Arg

465	470	475	480
Ala Ala Leu Glu Ser Gln Leu Arg Gln Ala Lys Thr Glu Leu Glu Glu	485	490	495
Thr Thr Ala Glu Ala Glu Glu Glu Ile Gln Ala Leu Thr Ala His Arg	500	505	510
Asp Glu Ile Gln Arg Lys Phe Asp Ala Leu Arg Asn Ser Cys Thr Val	515	520	525
Ile Thr Asp Leu Glu Glu Gln Leu Asn Gln Leu Thr Glu Asp Asn Ala	530	535	540
Glu Leu Asn Asn Gln Asn Phe Tyr Leu Ser Lys Gln Leu Asp Glu Ala	545	550	555
Ser Gly Ala Asn Asp Glu Ile Val Gln Leu Arg Ser Glu Val Asp His	565	570	575
Leu Arg Arg Glu Ile Thr Glu Arg Glu Met Gln Leu Thr Ser Gln Lys	580	585	590
Gln Thr Met Glu Ala Leu Lys Thr Thr Cys Thr Met Leu Glu Glu Gln	595	600	605
Val Leu Asp Leu Glu Ala Leu Asn Asp Glu Leu Leu Glu Lys Glu Arg	610	615	620
Gln Trp Glu Ala Trp Arg Ser Val Leu Gly Asp Glu Lys Ser Gln Phe	625	630	635
Glu Cys Arg Val Arg Glu Leu Gln Arg Met Leu Asp Thr Glu Lys Gln	645	650	655
Ser Arg Ala Arg Ala Asp Gln Arg Ile Thr Glu Ser Arg Gln Val Val	660	665	670
Glu Leu Ala Val Lys Glu His Lys Ala Glu Ile Leu Ala Leu Gln Gln	675	680	685
Ala Leu Lys Glu Gln Lys Leu Lys Ala Glu Ser Leu Ser Asp Lys Leu	690	695	700
Asn Asp Leu Glu Lys Lys His Ala Met Leu Glu Met Asn Ala Arg Ser	705	710	715
Leu Gln Gln Lys Leu Glu Thr Glu Arg Glu Leu Lys Gln Arg Leu Leu	725	730	735
Glu Glu Gln Ala Lys Leu Gln Gln Gln Met Asp Leu Gln Lys Asn His	740	745	750
Ile Phe Arg Leu Thr Gln Gly Leu Gln Glu Ala Leu Asp Arg Ala Asp	755	760	765
Leu Leu Lys Thr Glu Arg Ser Asp Leu Glu Tyr Gln Leu Glu Asn Ile			

770					775					780					
Gln	Val	Leu	Tyr	Ser	His	Glu	Lys	Val	Lys	Met	Glu	Gly	Thr	Ile	Ser
785					790					795					800
Gln	Gln	Thr	Lys	Leu	Ile	Asp	Phe	Leu	Gln	Ala	Lys	Met	Asp	Gln	Pro
			805						810					815	
Ala	Lys	Lys	Lys	Lys	Val	Pro	Leu	Gln	Tyr	Asn	Glu	Leu	Lys	Leu	Ala
			820					825					830		
Leu	Glu	Lys	Glu	Lys	Ala	Arg	Cys	Ala	Glu	Leu	Glu	Glu	Ala	Leu	Gln
	835					840						845			
Lys	Thr	Arg	Ile	Glu	Leu	Arg	Ser	Ala	Arg	Glu	Glu	Ala	Ala	His	Arg
	850					855					860				
Lys	Ala	Thr	Asp	His	Pro	His	Pro	Ser	Thr	Pro	Ala	Thr	Ala	Arg	Gln
865					870					875					880
Gln	Ile	Ala	Met	Ser	Ala	Ile	Val	Arg	Ser	Pro	Glu	His	Gln	Pro	Ser
			885						890					895	
Ala	Met	Ser	Leu	Leu	Ala	Pro	Pro	Ser	Ser	Arg	Arg	Lys	Glu	Ser	Ser
			900					905					910		
Thr	Pro	Glu	Glu	Phe	Ser	Arg	Arg	Leu	Lys	Glu	Arg	Met	His	His	Asn
	915						920					925			
Ile	Pro	His	Arg	Phe	Asn	Val	Gly	Leu	Asn	Met	Arg	Ala	Thr	Lys	Cys
	930					935					940				
Ala	Val	Cys	Leu	Asp	Thr	Val	His	Phe	Gly	Arg	Gln	Ala	Ser	Lys	Cys
945					950					955					960
Leu	Glu	Cys	Gln	Val	Met	Cys	His	Pro	Lys	Cys	Ser	Thr	Cys	Leu	Pro
			965						970					975	
Ala	Thr	Cys	Gly	Leu	Pro	Ala	Glu	Tyr	Ala	Thr	His	Phe	Thr	Glu	Ala
			980					985					990		
Phe	Cys	Arg	Asp	Lys	Met	Asn	Ser	Pro	Gly	Leu	Gln	Ser	Lys	Glu	Pro
	995					1000					1005				
Gly	Ser	Ser	Leu	His	Leu	Glu	Gly	Trp	Met	Lys	Val	Pro	Arg	Asn	Asn
	1010					1015					1020				
Lys	Arg	Gly	Gln	Gln	Gly	Trp	Asp	Arg	Lys	Tyr	Ile	Val	Leu	Glu	Gly
1025					1030					1035					1040
Ser	Lys	Val	Leu	Ile	Tyr	Asp	Asn	Glu	Ala	Arg	Glu	Ala	Gly	Gln	Arg
			1045					1050					1055		
Pro	Val	Glu	Glu	Phe	Glu	Leu	Cys	Leu	Pro	Asp	Gly	Asp	Val	Ser	Ile
			1060					1065					1070		
His	Gly	Ala	Val	Gly	Ala	Ser	Glu	Leu	Ala	Asn	Thr	Ala	Lys	Ala	Asp

1075	1080	1085
Val Pro Tyr Ile Leu Lys Met Glu Ser His Pro His Thr Thr Cys Trp 1090	1095	1100
Pro Gly Arg Thr Leu Tyr Leu Leu Ala Pro Ser Phe Pro Asp Lys Gln 1105	1110	1115 1120
Arg Trp Val Thr Ala Leu Glu Ser Val Val Ala Gly Gly Arg Val Ser 1125	1130	1135
Arg Glu Lys Ala Glu Ala Asp Ala Lys Leu Leu Gly Asn Ser Leu Leu 1140	1145	1150
Lys Leu Glu Gly Asp Asp Arg Leu Asp Met Asn Cys Thr Leu Pro Phe 1155	1160	1165
Ser Asp Gln Val Val Leu Val Gly Thr Glu Glu Gly Leu Tyr Ala Leu 1170	1175	1180
Asn Val Leu Lys Asn Ser Leu Thr His Ile Pro Gly Ile Gly Ala Val 1185	1190	1195 1200
Phe Gln Ile Tyr Ile Ile Lys Asp Leu Glu Lys Leu Leu Met Ile Ala 1205	1210	1215
Gly Glu Glu Arg Ala Leu Cys Leu Val Asp Val Lys Lys Val Lys Gln 1220	1225	1230
Ser Leu Ala Gln Ser His Leu Pro Ala Gln Pro Asp Val Ser Pro Asn 1235	1240	1245
Ile Phe Glu Ala Val Lys Gly Cys His Leu Phe Ala Ala Gly Lys Ile 1250	1255	1260
Glu Asn Ser Leu Cys Ile Cys Ala Ala Met Pro Ser Lys Val Val Ile 1265	1270	1275 1280
Leu Arg Tyr Asn Asp Asn Leu Ser Lys Tyr Cys Ile Arg Lys Glu Ile 1285	1290	1295
Glu Thr Ser Glu Pro Cys Ser Cys Ile His Phe Thr Asn Tyr Ser Ile 1300	1305	1310
Leu Ile Gly Thr Asn Lys Phe Tyr Glu Ile Asp Met Lys Gln Tyr Thr 1315	1320	1325
Leu Asp Glu Phe Leu Asp Lys Asn Asp His Ser Leu Ala Pro Ala Val 1330	1335	1340
Phe Ala Ser Ser Ser Asn Ser Phe Pro Val Ser Ile Val Gln Ala Asn 1345	1350	1355 1360
Ser Ala Gly Gln Arg Glu Glu Tyr Leu Leu Cys Phe His Glu Phe Gly 1365	1370	1375
Val Phe Val Asp Ser Tyr Gly Arg Arg Ser Arg Thr Asp Asp Leu Lys		

1380	1385	1390
Trp Ser Arg Leu Pro Leu Ala Phe Ala Tyr Arg Glu Pro Tyr Leu Phe 1395 1400 1405		
Val Thr His Phe Asn Ser Leu Glu Val Ile Glu Ile Gln Ala Arg Ser 1410 1415 1420		
Ser Leu Gly Ser Pro Ala Arg Ala Tyr Leu Glu Ile Pro Asn Pro Arg 1425 1430 1435 1440		
Tyr Leu Gly Pro Ala Ile Ser Ser Gly Ala Ile Tyr Leu Ala Ser Ser 1445 1450 1455		
Tyr Gln Asp Lys Leu Arg Val Ile Cys Cys Lys Gly Asn Leu Val Lys 1460 1465 1470		
Glu Ser Gly Thr Glu Gln His Arg Val Pro Ser Thr Ser Arg Ser Ser 1475 1480 1485		
Pro Asn Lys Arg Gly Pro Pro Thr Tyr Asn Glu His Ile Thr Lys Arg 1490 1495 1500		
Val Ala Ser Ser Pro Ala Pro Pro Glu Gly Pro Ser His Pro Arg Glu 1505 1510 1515 1520		
Pro Ser Thr Pro His Arg Tyr Arg Asp Arg Glu Gly Arg Thr Glu Leu 1525 1530 1535		
Arg Arg Asp Lys Ser Pro Gly Arg Pro Leu Glu Arg Glu Lys Ser Pro 1540 1545 1550		
Gly Arg Met Leu Ser Thr Arg Arg Glu Arg Ser Pro Gly Arg Leu Phe 1555 1560 1565		
Glu Asp Ser Ser Arg Gly Arg Leu Pro Ala Gly Ala Val Arg Thr Pro 1570 1575 1580		
Leu Ser Gln Val Asn Lys Val Trp Asp Gln Ser Ser Val 1585 1590 1595		
<210> 42		
<211> 1871		
<212> PRT		
<213> Homo sapiens		
<400> 42		
Met Pro Ser Val Cys Leu Leu Leu Leu Leu Phe Leu Ala Val Gly Gly 1 5 10 15		
Ala Leu Gly Asn Arg Pro Phe Arg Ala Phe Val Val Thr Asp Thr Thr 20 25 30		
Leu Thr His Leu Ala Val His Arg Val Thr Gly Glu Val Phe Val Gly 35 40 45		

Ala Val Asn Arg Val Phe Lys Leu Ala Pro Asn Leu Thr Glu Leu Arg
 50 55 60
 Ala His Val Thr Gly Pro Val Glu Asp Asn Ala Arg Cys Tyr Pro Pro
 65 70 75 80
 Pro Ser Met Arg Val Cys Ala His Arg Leu Ala Pro Val Asp Asn Ile
 85 90 95
 Asn Lys Leu Leu Leu Ile Asp Tyr Ala Ala Arg Arg Leu Val Ala Cys
 100 105 110
 Gly Ser Ile Trp Gln Gly Ile Cys Gln Phe Leu Arg Leu Asp Asp Leu
 115 120 125
 Phe Lys Leu Gly Glu Pro His His Arg Lys Glu His Tyr Leu Ser Gly
 130 135 140
 Ala Gln Glu Pro Asp Ser Met Ala Gly Val Ile Val Glu Gln Gly Gln
 145 150 155 160
 Gly Pro Ser Lys Leu Phe Val Gly Thr Ala Val Asp Gly Lys Ser Glu
 165 170 175
 Tyr Phe Pro Thr Leu Ser Ser Arg Lys Leu Ile Ser Asp Glu Asp Ser
 180 185 190
 Ala Asp Met Phe Ser Leu Val Tyr Gln Asp Glu Phe Val Ser Ser Gln
 195 200 205
 Ile Lys Ile Pro Ser Asp Thr Leu Ser Leu Tyr Pro Ala Phe Asp Ile
 210 215 220
 Tyr Tyr Ile Tyr Gly Phe Val Ser Ala Ser Phe Val Tyr Phe Leu Thr
 225 230 235 240
 Leu Gln Leu Asp Thr Gln Gln Thr Leu Leu Asp Thr Ala Gly Glu Lys
 245 250 255
 Phe Phe Thr Ser Lys Ile Val Arg Met Cys Ala Gly Asp Ser Glu Phe
 260 265 270
 Tyr Ser Tyr Val Glu Phe Pro Ile Gly Cys Ser Trp Arg Gly Val Glu
 275 280 285
 Tyr Arg Leu Val Gln Ser Ala His Leu Ala Lys Pro Gly Leu Leu Leu
 290 295 300
 Ala Gln Ala Leu Gly Val Pro Ala Asp Glu Asp Val Leu Phe Thr Ile
 305 310 315 320
 Phe Ser Gln Gly Gln Lys Asn Arg Ala Ser Pro Pro Arg Gln Thr Ile
 325 330 335
 Leu Cys Leu Phe Thr Leu Ser Asn Ile Asn Ala His Ile Arg Arg Arg
 340 345 350

Ile Gln Ser Cys Tyr Arg Gly Glu Gly Thr Leu Ala Leu Pro Trp Leu
 355 360 365
 Leu Asn Lys Glu Leu Pro Cys Ile Asn Thr Pro Met Gln Ile Asn Gly
 370 375 380
 Asn Phe Cys Gly Leu Val Leu Asn Gln Pro Leu Gly Gly Leu His Val
 385 390 395 400
 Ile Glu Gly Leu Pro Leu Leu Ala Asp Ser Thr Asp Gly Met Ala Ser
 405 410 415
 Val Ala Ala Tyr Thr Tyr Arg Gln His Ser Val Val Phe Ile Gly Thr
 420 425 430
 Arg Ser Gly Ser Leu Lys Lys Val Arg Val Asp Gly Phe Gln Asp Ala
 435 440 445
 His Leu Tyr Glu Thr Val Pro Val Val Asp Gly Ser Pro Ile Leu Arg
 450 455 460
 Asp Leu Leu Phe Ser Pro Asp His Arg His Ile Tyr Leu Leu Ser Glu
 465 470 475 480
 Lys Gln Val Ser Gln Leu Pro Val Glu Thr Cys Glu Gln Tyr Gln Ser
 485 490 495
 Cys Ala Ala Cys Leu Gly Ser Gly Asp Pro His Cys Gly Trp Cys Val
 500 505 510
 Leu Arg His Arg Cys Cys Arg Glu Gly Ala Cys Leu Gly Ala Ser Ala
 515 520 525
 Pro His Gly Phe Ala Glu Glu Leu Ser Lys Cys Val Gln Val Arg Val
 530 535 540
 Arg Pro Asn Asn Val Ser Val Thr Ser Pro Gly Val Gln Leu Thr Val
 545 550 555 560
 Thr Leu His Asn Val Pro Asp Leu Ser Ala Gly Val Ser Cys Ala Phe
 565 570 575
 Glu Ala Ala Ala Glu Asn Glu Ala Val Leu Leu Pro Ser Gly Glu Leu
 580 585 590
 Leu Cys Pro Ser Pro Ser Leu Gln Glu Leu Arg Ala Leu Thr Arg Gly
 595 600 605
 His Gly Ala Thr Arg Thr Val Arg Leu Gln Leu Leu Ser Lys Glu Thr
 610 615 620
 Gly Val Arg Phe Ala Gly Ala Asp Phe Val Phe Tyr Asn Cys Ser Val
 625 630 635 640
 Leu Gln Ser Cys Met Ser Cys Val Gly Ser Pro Tyr Pro Cys His Trp
 645 650 655

Cys Lys Tyr Arg His Thr Cys Thr Ser Arg Pro His Glu Cys Ser Phe
 660 665 670
 Gln Glu Gly Arg Val His Ser Pro Glu Gly Cys Pro Glu Ile Leu Pro
 675 680 685
 Ser Gly Asp Leu Leu Ile Pro Val Gly Val Met Gln Pro Leu Thr Leu
 690 695 700
 Arg Ala Lys Asn Leu Pro Gln Pro Gln Ser Gly Gln Lys Asn Tyr Glu
 705 710 715 720
 Cys Val Val Arg Val Gln Gly Arg Gln Gln Arg Val Pro Ala Val Arg
 725 730 735
 Phe Asn Ser Ser Ser Val Gln Cys Gln Asn Ala Ser Tyr Ser Tyr Glu
 740 745 750
 Gly Asp Glu His Gly Asp Thr Glu Leu Asp Phe Ser Val Val Trp Asp
 755 760 765
 Gly Asp Phe Pro Ile Asp Lys Pro Pro Ser Phe Arg Ala Leu Leu Tyr
 770 775 780
 Lys Cys Trp Ala Gln Arg Pro Ser Cys Gly Leu Cys Leu Lys Ala Asp
 785 790 795 800
 Pro Arg Phe Asn Cys Gly Trp Cys Ile Ser Glu His Arg Cys Gln Leu
 805 810 815
 Arg Thr His Cys Pro Ala Pro Lys Thr Asn Trp Met His Leu Ser Gln
 820 825 830
 Lys Gly Thr Arg Cys Ser His Pro Arg Ile Thr Gln Ile His Pro Leu
 835 840 845
 Val Gly Pro Lys Glu Gly Gly Thr Arg Val Thr Ile Val Gly Asp Asn
 850 855 860
 Leu Gly Leu Leu Ser Arg Glu Val Gly Leu Arg Val Ala Gly Val Arg
 865 870 875 880
 Cys Asn Ser Ile Pro Ala Glu Tyr Ile Ser Ala Glu Arg Ile Val Cys
 885 890 895
 Glu Met Glu Glu Ser Leu Val Pro Ser Pro Pro Pro Gly Pro Val Glu
 900 905 910
 Leu Cys Val Gly Asp Cys Ser Ala Asp Phe Arg Thr Gln Ser Glu Gln
 915 920 925
 Val Tyr Ser Phe Val Thr Pro Thr Phe Asp Gln Val Ser Pro Ser Arg
 930 935 940
 Gly Pro Ala Ser Gly Gly Thr Arg Leu Thr Ile Ser Gly Ser Ser Leu
 945 950 955 960

Asp Ala Gly Ser Arg Val Thr Val Thr Val Arg Asp Ser Glu Cys Gln
 965 970 975
 Phe Val Arg Arg Asp Ala Lys Ala Ile Val Cys Ile Ser Pro Leu Ser
 980 985 990
 Thr Leu Gly Pro Ser Gln Ala Pro Ile Thr Leu Ala Ile Asp Arg Ala
 995 1000 1005
 Asn Ile Ser Ser Pro Gly Leu Ile Tyr Thr Tyr Thr Gln Asp Pro Thr
 1010 1015 1020
 Val Thr Arg Leu Glu Pro Thr Trp Ser Ile Ile Asn Gly Ser Thr Ala
 1025 1030 1035 1040
 Ile Thr Val Ser Gly Thr His Leu Leu Thr Val Gln Glu Pro Arg Val
 1045 1050 1055
 Arg Ala Lys Tyr Arg Gly Ile Glu Thr Thr Asn Thr Cys Gln Val Ile
 1060 1065 1070
 Asn Asp Thr Ala Met Leu Cys Lys Ala Pro Gly Ile Phe Leu Gly Arg
 1075 1080 1085
 Pro Gln Pro Arg Ala Gln Gly Glu His Pro Asp Glu Phe Gly Phe Leu
 1090 1095 1100
 Leu Asp His Val Gln Thr Ala Arg Ser Leu Asn Arg Ser Ser Phe Thr
 1105 1110 1115 1120
 Tyr Tyr Pro Asp Pro Ser Phe Glu Pro Leu Gly Pro Ser Gly Val Leu
 1125 1130 1135
 Asp Val Lys Pro Gly Ser His Val Val Leu Lys Gly Lys Asn Leu Ile
 1140 1145 1150
 Pro Ala Ala Ala Gly Ser Ser Arg Leu Asn Tyr Thr Val Leu Ile Gly
 1155 1160 1165
 Gly Gln Pro Cys Ser Leu Thr Val Ser Asp Thr Gln Leu Leu Cys Asp
 1170 1175 1180
 Ser Pro Ser Gln Thr Gly Arg Gln Pro Val Met Val Leu Val Gly Gly
 1185 1190 1195 1200
 Leu Glu Phe Trp Leu Gly Thr Leu His Ile Ser Ala Glu Arg Ala Leu
 1205 1210 1215
 Thr Leu Pro Ala Met Met Gly Leu Ala Ala Gly Gly Gly Leu Leu Leu
 1220 1225 1230
 Leu Ala Ile Thr Ala Val Leu Val Ala Tyr Lys Arg Lys Thr Gln Asp
 1235 1240 1245
 Ala Asp Arg Thr Leu Lys Arg Leu Gln Leu Gln Met Asp Asn Leu Glu
 1250 1255 1260

Ser Arg Val Ala Leu Glu Cys Lys Glu Ala Phe Ala Glu Leu Gln Thr
 1265 1270 1275 1280

Asp Ile Asn Glu Leu Thr Asn His Met Asp Glu Val Gln Ile Pro Phe
 1285 1290 1295

Leu Asp Tyr Arg Thr Tyr Ala Val Arg Val Leu Phe Pro Gly Ile Glu
 1300 1305 1310

Ala His Pro Val Leu Lys Glu Leu Asp Thr Pro Pro Asn Val Glu Lys
 1315 1320 1325

Ala Leu Arg Leu Phe Gly Gln Leu Leu His Ser Arg Ala Phe Val Leu
 1330 1335 1340

Thr Phe Ile His Thr Leu Glu Ala Gln Ser Ser Phe Ser Met Arg Asp
 1345 1350 1355 1360

Arg Gly Thr Val Ala Ser Leu Thr Met Val Ala Leu Gln Ser Arg Leu
 1365 1370 1375

Asp Tyr Ala Thr Gly Leu Leu Lys Gln Leu Leu Ala Asp Leu Ile Glu
 1380 1385 1390

Lys Asn Leu Glu Ser Lys Asn His Pro Lys Leu Leu Leu Arg Arg Thr
 1395 1400 1405

Glu Ser Val Ala Glu Lys Met Leu Thr Asn Trp Phe Thr Phe Leu Leu
 1410 1415 1420

His Lys Phe Leu Lys Glu Cys Ala Gly Glu Pro Leu Phe Leu Leu Tyr
 1425 1430 1435 1440

Cys Ala Ile Lys Gln Gln Met Glu Lys Gly Pro Ile Asp Ala Ile Thr
 1445 1450 1455

Gly Glu Ala Arg Tyr Ser Leu Ser Glu Asp Lys Leu Ile Arg Gln Gln
 1460 1465 1470

Ile Asp Tyr Lys Thr Leu Thr Leu His Cys Val Cys Pro Glu Asn Glu
 1475 1480 1485

Gly Ser Ala Gln Val Pro Val Lys Val Leu Asn Cys Asp Ser Ile Thr
 1490 1495 1500

Gln Ala Lys Asp Lys Leu Leu Asp Thr Val Tyr Lys Gly Ile Pro Tyr
 1505 1510 1515 1520

Ser Gln Arg Pro Lys Ala Glu Asp Met Asp Leu Glu Trp Arg Gln Gly
 1525 1530 1535

Arg Met Thr Arg Ile Ile Leu Gln Asp Glu Asp Val Thr Thr Lys Ile
 1540 1545 1550

Glu Cys Asp Trp Lys Arg Leu Asn Ser Leu Ala His Tyr Gln Val Thr
 1555 1560 1565

Asp Gly Ser Leu Val Ala Leu Val Pro Lys Gln Val Ser Ala Tyr Asn
 1570 1575 1580
 Met Ala Asn Ser Phe Thr Phe Thr Arg Ser Leu Ser Arg Tyr Glu Ser
 1585 1590 1595 1600
 Leu Leu Arg Thr Ala Ser Ser Pro Asp Ser Leu Arg Ser Arg Ala Pro
 1605 1610 1615
 Met Ile Thr Pro Asp Gln Glu Thr Gly Thr Lys Leu Trp His Leu Val
 1620 1625 1630
 Lys Asn His Asp His Ala Asp His Arg Glu Gly Asp Arg Gly Ser Lys
 1635 1640 1645
 Met Val Ser Glu Ile Tyr Leu Thr Arg Leu Leu Ala Thr Lys Gly Thr
 1650 1655 1660
 Leu Gln Lys Phe Val Asp Asp Leu Phe Glu Thr Val Phe Ser Thr Ala
 1665 1670 1675 1680
 His Arg Gly Ser Ala Leu Pro Leu Ala Ile Lys Tyr Met Phe Asp Phe
 1685 1690 1695
 Leu Asp Glu Gln Ala Asp Gln Arg Gln Ile Ser Asp Pro Asp Val Arg
 1700 1705 1710
 His Thr Trp Lys Ser Asn Cys Leu Pro Leu Arg Phe Trp Val Asn Val
 1715 1720 1725
 Ile Lys Asn Pro Gln Phe Val Phe Asp Ile His Lys Asn Ser Ile Thr
 1730 1735 1740
 Asp Ala Cys Leu Ser Val Val Ala Gln Thr Phe Met Asp Ser Cys Ser
 1745 1750 1755 1760
 Thr Ser Glu His Arg Leu Gly Lys Asp Ser Pro Ser Asn Lys Leu Leu
 1765 1770 1775
 Tyr Ala Lys Asp Ile Pro Asn Tyr Lys Ser Trp Val Glu Arg Tyr Tyr
 1780 1785 1790
 Arg Asp Ile Ala Lys Met Ala Ser Ile Ser Asp Gln Asp Met Asp Ala
 1795 1800 1805
 Tyr Leu Val Glu Gln Ser Arg Leu His Ala Ser Asp Phe Ser Val Leu
 1810 1815 1820
 Ser Ala Leu Asn Glu Leu Tyr Phe Tyr Val Thr Lys Tyr Arg Gln Glu
 1825 1830 1835 1840
 Ile Leu Thr Ala Leu Asp Arg Asp Ala Ser Cys Arg Lys His Lys Leu
 1845 1850 1855
 Arg Gln Lys Leu Glu Gln Ile Ile Ser Leu Val Ser Ser Asp Ser
 1860 1865 1870

<400> 43

Ser Arg Arg Pro Gln Gln Pro Ala Ala Ala Glu Val Asp Arg Ala Gly
20 25 30

Thr Glu Gly Gln Thr Asp Val Ala Glu Leu Glu Ser Cys Glu Gly Gln
35 40 45

Pro Gly Lys Val Glu Gln Met Ser Thr His Arg Ser Arg Leu Leu Thr
50 55 60

Ala Ala Pro Leu Ser Met Glu Gln Arg Arg Pro Trp Pro Arg Ala Leu
65 70 75 80

Glu Val Asp Ser Arg Ser Val Val Leu Leu Ser Val Val Trp Val Leu
85 90 95

Leu Ala Pro Pro Ala Ala Gly Met Pro Gln Phe Ser Thr Phe His Ser
100 105 110

Glu Asn Arg Asp Trp Thr Phe Asn His Leu Thr Val His Gln Gly Thr
115 120 125

Gly Ala Val Tyr Val Gly Ala Ile Asn Arg Val Tyr Lys Leu Thr Gly
130 135 140

Asn Leu Thr Ile Gln Val Ala His Lys Thr Gly Pro Glu Glu Asp Asn
145 150 155 160

Lys Ser Cys Tyr Pro Pro Leu Ile Val Gln Pro Cys Ser Glu Val Leu
165 170 175

Thr Leu Thr Asn Asn Val Asn Lys Leu Leu Ile Ile Asp Tyr Ser Glu
180 185 190

Asn Arg Leu Leu Ala Cys Gly Ser Leu Tyr Gln Gly Val Cys Lys Leu
195 200 205

Leu Arg Leu Asp Asp Leu Phe Ile Leu Val Glu Pro Ser His Lys Lys
210 215 220

Glu His Tyr Leu Ser Ser Val Asn Lys Thr Gly Thr Met Tyr Gly Val
225 230 235 240

Ile Val Arg Ser Glu Gly Glu Asp Gly Lys Leu Phe Ile Gly Thr Ala
245 250 255

Val Asp Gly Lys Gln Asp Tyr Phe Pro Thr Leu Ser Ser Arg Lys Leu
260 265 270

Pro Arg Asp Pro Glu Ser Ser Ala Met Leu Asp Tyr Glu Leu His Ser
 275 280 285
 Asp Phe Val Ser Ser Leu Ile Lys Ile Pro Ser Asp Thr Leu Ala Leu
 290 295 300
 Val Ser His Phe Asp Ile Phe Tyr Ile Tyr Gly Phe Ala Ser Gly Gly
 305 310 315 320
 Phe Val Tyr Phe Leu Thr Val Gln Pro Glu Thr Pro Glu Gly Val Ala
 325 330 335
 Ile Asn Ser Ala Gly Asp Leu Phe Tyr Thr Ser Arg Ile Val Arg Leu
 340 345 350
 Cys Lys Asp Asp Pro Lys Phe His Ser Tyr Val Ser Leu Pro Phe Gly
 355 360 365
 Cys Thr Arg Ala Gly Val Glu Tyr Arg Leu Leu Gln Ala Ala Tyr Leu
 370 375 380
 Ala Lys Pro Gly Asp Ser Leu Ala Gln Ala Phe Asn Ile Thr Ser Gln
 385 390 395 400
 Asp Asp Val Leu Phe Ala Ile Phe Ser Lys Gly Gln Lys Gln Tyr His
 405 410 415
 His Pro Pro Asp Asp Ser Ala Leu Cys Ala Phe Pro Ile Arg Ala Ile
 420 425 430
 Asn Leu Gln Ile Lys Gly Arg Leu Gln Ser Cys Tyr Gln Gly Glu Gly
 435 440 445
 Asn Leu Glu Leu Asn Trp Leu Leu Gly Lys Asp Val Gln Cys Thr Lys
 450 455 460
 Ala Pro Val Pro Ile Asp Asp Asn Phe Cys Gly Leu Asp Ile Asn Gln
 465 470 475 480
 Pro Leu Gly Gly Ser Thr Pro Val Glu Gly Leu Thr Leu Tyr Thr Thr
 485 490 495
 Ser Arg Asp Arg Met Thr Ser Val Ala Ser Tyr Val Tyr Asn Gly Tyr
 500 505 510
 Ser Val Val Phe Val Gly Thr Lys Ser Gly Lys Leu Lys Lys Ile Arg
 515 520 525
 Ala Asp Gly Pro Pro His Gly Gly Val Gln Tyr Glu Met Val Ser Val
 530 535 540
 Leu Lys Asp Gly Ser Pro Ile Leu Arg Asp Met Ala Phe Ser Ile Asp
 545 550 555 560
 Gln Arg Tyr Leu Tyr Val Met Ser Glu Arg Gln Val Thr Arg Val Pro
 565 570 575

Val	Glu	Ser	Cys	Glu	Gln	Tyr	Thr	Thr	Cys	Gly	Glu	Cys	Leu	Ser	Ser	580	585	590
Gly	Asp	Pro	His	Cys	Gly	Trp	Cys	Ala	Leu	His	Asn	Met	Cys	Ser	Arg	595	600	605
Arg	Asp	Lys	Cys	Gln	Gln	Ala	Trp	Glu	Pro	Asn	Arg	Phe	Ala	Ala	Ser	610	615	620
Ile	Ser	Gln	Cys	Val	Ser	Leu	Ala	Val	His	Pro	Ser	Ser	Ile	Ser	Val	625	630	635
Ser	Glu	His	Ser	Arg	Leu	Leu	Ser	Leu	Val	Val	Ser	Asp	Ala	Pro	Asp	645	650	655
Leu	Ser	Ala	Gly	Ile	Ala	Cys	Ala	Phe	Gly	Asn	Leu	Thr	Glu	Val	Glu	660	665	670
Gly	Gln	Val	Ser	Gly	Ser	Gln	Val	Ile	Cys	Ile	Ser	Pro	Gly	Pro	Lys	675	680	685
Asp	Val	Pro	Val	Ile	Pro	Leu	Asp	Gln	Asp	Trp	Phe	Gly	Leu	Glu	Leu	690	695	700
Gln	Leu	Arg	Ser	Lys	Glu	Thr	Gly	Lys	Ile	Phe	Val	Ser	Thr	Glu	Phe	705	710	715
Lys	Phe	Tyr	Asn	Cys	Ser	Ala	His	Gln	Leu	Cys	Leu	Ser	Cys	Val	Asn	725	730	735
Ser	Ala	Phe	Arg	Cys	His	Trp	Cys	Lys	Tyr	Arg	Asn	Leu	Cys	Thr	His	740	745	750
Asp	Pro	Thr	Thr	Cys	Ser	Phe	Gln	Glu	Gly	Arg	Ile	Asn	Ile	Ser	Glu	755	760	765
Asp	Cys	Pro	Gln	Leu	Val	Pro	Thr	Glu	Glu	Ile	Leu	Ile	Pro	Val	Gly	770	775	780
Glu	Val	Lys	Pro	Ile	Thr	Leu	Lys	Ala	Arg	Asn	Leu	Pro	Gln	Pro	Gln	785	790	795
Ser	Gly	Gln	Arg	Gly	Tyr	Glu	Cys	Val	Leu	Asn	Ile	Gln	Gly	Ala	Ile	805	810	815
His	Arg	Val	Pro	Ala	Leu	Arg	Phe	Asn	Ser	Ser	Ser	Val	Gln	Cys	Gln	820	825	830
Asn	Ser	Ser	Tyr	Gln	Tyr	Asp	Gly	Met	Asp	Ile	Ser	Asn	Leu	Ala	Val	835	840	845
Asp	Phe	Ala	Val	Val	Trp	Asn	Gly	Asn	Phe	Ile	Ile	Asp	Asn	Pro	Gln	850	855	860
Asp	Leu	Lys	Val	His	Leu	Tyr	Lys	Cys	Ala	Ala	Gln	Arg	Glu	Ser	Cys	865	870	875

Gly Leu Cys Leu Lys Ala Asp Arg Lys Phe Glu Cys Gly Trp Cys Ser
 885 890 895
 Gly Glu Arg Arg Cys Thr Leu His Gln His Cys Thr Ser Pro Ser Ser
 900 905 910
 Pro Trp Leu Asp Trp Ser Ser His Asn Val Lys Cys Ser Asn Pro Gln
 915 920 925
 Ile Thr Glu Ile Leu Thr Val Ser Gly Pro Pro Glu Gly Gly Thr Arg
 930 935 940
 Val Thr Ile His Gly Val Asn Leu Gly Leu Asp Phe Ser Glu Ile Ala
 945 950 955 960
 His His Val Gln Val Ala Gly Val Pro Cys Thr Pro Leu Pro Gly Glu
 965 970 975
 Tyr Ile Ile Ala Glu Gln Ile Val Cys Glu Met Gly His Ala Leu Val
 980 985 990
 Gly Thr Thr Ser Gly Pro Val Arg Leu Cys Ile Gly Glu Cys Lys Pro
 995 1000 1005
 Glu Phe Met Thr Lys Ser His Gln Gln Tyr Thr Phe Val Asn Pro Ser
 1010 1015 1020
 Val Leu Ser Leu Asn Pro Ile Arg Gly Pro Glu Ser Gly Gly Thr Met
 1025 1030 1035 1040
 Val Thr Ile Thr Gly His Tyr Leu Gly Ala Gly Ser Ser Val Ala Val
 1045 1050 1055
 Tyr Leu Gly Asn Gln Thr Cys Glu Phe Tyr Gly Arg Ser Met Ser Glu
 1060 1065 1070
 Ile Val Cys Val Ser Pro Pro Ser Ser Asn Gly Leu Gly Pro Val Pro
 1075 1080 1085
 Val Ser Val Ser Val Asp Arg Ala His Val Asp Ser Asn Leu Gln Phe
 1090 1095 1100
 Glu Tyr Ile Asp Asp Pro Arg Val Gln Arg Ile Glu Pro Glu Trp Ser
 1105 1110 1115 1120
 Ile Ala Ser Gly His Thr Pro Leu Thr Ile Thr Gly Phe Asn Leu Asp
 1125 1130 1135
 Val Ile Gln Glu Pro Arg Ile Arg Val Lys Phe Asn Gly Lys Glu Ser
 1140 1145 1150
 Val Asn Val Cys Lys Val Val Asn Thr Thr Thr Leu Thr Cys Leu Ala
 1155 1160 1165
 Pro Ser Leu Thr Thr Asp Tyr Arg Pro Gly Leu Asp Thr Val Glu Arg
 1170 1175 1180

Pro Asp Glu Phe Gly Phe Val Phe Asn Asn Val Gln Ser Leu Leu Ile
 1185 1190 1195 1200
 Tyr Asn Asp Thr Lys Phe Ile Tyr Tyr Pro Asn Pro Thr Phe Glu Leu
 1205 1210 1215
 Leu Ser Pro Thr Gly Val Leu Asp Gln Lys Pro Gly Ser Pro Ile Ile
 1220 1225 1230
 Leu Lys Gly Lys Asn Leu Cys Pro Pro Ala Ser Gly Gly Ala Lys Leu
 1235 1240 1245
 Asn Tyr Thr Val Leu Ile Gly Glu Thr Pro Cys Ala Val Thr Val Ser
 1250 1255 1260
 Glu Thr Gln Leu Leu Cys Glu Pro Pro Asn Leu Thr Gly Gln His Lys
 1265 1270 1275 1280
 Val Met Val His Val Gly Gly Met Val Phe Ser Pro Gly Ser Val Ser
 1285 1290 1295
 Val Ile Ser Asp Ser Leu Leu Thr Leu Pro Ala Ile Val Ser Ile Ala
 1300 1305 1310
 Ala Gly Gly Ser Leu Leu Leu Ile Ile Val Ile Ile Val Leu Ile Ala
 1315 1320 1325
 Tyr Lys Arg Lys Ser Arg Glu Asn Asp Leu Thr Leu Lys Arg Leu Gln
 1330 1335 1340
 Met Gln Met Asp Asn Leu Glu Ser Arg Val Ala Leu Glu Cys Lys Glu
 1345 1350 1355 1360
 Ala Phe Ala Glu Leu Gln Thr Asp Ile Asn Glu Leu Thr Ser Asp Leu
 1365 1370 1375
 Asp Arg Ser Gly Ile Pro Tyr Leu Asp Tyr Arg Thr Tyr Ala Met Arg
 1380 1385 1390
 Val Leu Phe Pro Gly Ile Glu Asp His Pro Val Leu Arg Glu Leu Glu
 1395 1400 1405
 Val Gln Gly Asn Gly Gln Gln His Val Glu Lys Ala Leu Lys Leu Phe
 1410 1415 1420
 Ala Gln Leu Ile Asn Asn Lys Val Phe Leu Leu Thr Phe Ile Arg Thr
 1425 1430 1435 1440
 Leu Glu Leu Gln Arg Ser Phe Ser Met Arg Asp Arg Gly Asn Val Ala
 1445 1450 1455
 Ser Leu Ile Met Thr Gly Leu Gln Gly Arg Leu Glu Tyr Ala Thr Asp
 1460 1465 1470
 Val Leu Lys Gln Leu Leu Ser Asp Leu Ile Asp Lys Asn Leu Glu Asn
 1475 1480 1485

Lys Asn His Pro Lys Leu Leu Leu Arg Arg Thr Glu Ser Val Ala Glu
 1490 1495 1500
 Lys Met Leu Thr Asn Trp Phe Ala Phe Leu Leu His Lys Phe Leu Lys
 1505 1510 1515 1520
 Glu Cys Ala Gly Glu Pro Leu Phe Met Leu Tyr Cys Ala Ile Lys Gln
 1525 1530 1535
 Gln Met Glu Lys Gly Pro Ile Asp Ala Ile Thr Gly Glu Ala Arg Tyr
 1540 1545 1550
 Ser Leu Ser Glu Asp Lys Leu Ile Arg Gln Gln Ile Glu Tyr Lys Thr
 1555 1560 1565
 Leu Ile Leu Asn Cys Val Asn Pro Asp Asn Glu Asn Ser Pro Glu Ile
 1570 1575 1580
 Pro Val Lys Val Leu Asn Cys Asp Thr Ile Thr Gln Val Lys Glu Lys
 1585 1590 1595 1600
 Ile Leu Asp Ala Val Tyr Lys Asn Val Pro Tyr Ser Gln Arg Pro Arg
 1605 1610 1615
 Ala Val Asp Met Asp Leu Glu Trp Arg Gln Gly Arg Ile Ala Arg Val
 1620 1625 1630
 Val Leu Gln Asp Glu Asp Ile Thr Thr Lys Ile Glu Gly Asp Trp Lys
 1635 1640 1645
 Arg Leu Asn Thr Leu Met His Tyr Gln Val Ser Asp Arg Ser Val Val
 1650 1655 1660
 Ala Leu Val Pro Lys Gln Thr Ser Ser Tyr Asn Ile Pro Ala Ser Ala
 1665 1670 1675 1680
 Ser Ile Ser Arg Thr Ser Ile Ser Arg Tyr Asp Ser Ser Phe Arg Tyr
 1685 1690 1695
 Thr Gly Ser Pro Asp Ser Leu Arg Ser Arg Ala Pro Met Ile Thr Pro
 1700 1705 1710
 Asp Leu Glu Ser Gly Val Lys Val Trp His Leu Val Lys Asn His Asp
 1715 1720 1725
 His Gly Asp Gln Lys Glu Gly Asp Arg Gly Ser Lys Met Val Ser Glu
 1730 1735 1740
 Ile Tyr Leu Thr Arg Leu Leu Ala Thr Lys Gly Thr Leu Gln Lys Phe
 1745 1750 1755 1760
 Val Asp Asp Leu Phe Glu Thr Leu Phe Ser Thr Val His Arg Gly Ser
 1765 1770 1775
 Ala Leu Pro Leu Ala Ile Lys Tyr Met Phe Asp Phe Leu Asp Glu Gln
 1780 1785 1790

Ala Asp Arg His Ser Ile His Asp Thr Asp Val Arg His Thr Trp Lys
 1795 1800 1805
 Ser Asn Cys Leu Pro Leu Arg Phe Trp Val Asn Val Ile Lys Asn Pro
 1810 1815 1820
 Gln Phe Val Phe Asp Ile His Lys Gly Ser Ile Thr Asp Ala Cys Leu
 1825 1830 1835 1840
 Ser Val Val Ala Gln Thr Phe Met Asp Ser Cys Ser Thr Ser Glu His
 1845 1850 1855
 Arg Leu Gly Lys Asp Ser Pro Ser Asn Lys Leu Leu Tyr Ala Lys Asp
 1860 1865 1870
 Ile Pro Ser Tyr Lys Ser Trp Val Glu Arg Tyr Tyr Ala Asp Ile Ala
 1875 1880 1885
 Lys Leu Pro Ala Ile Ser Asp Gln Asp Met Asn Ala Tyr Leu Ala Glu
 1890 1895 1900
 Gln Ser Arg Leu His Ala Val Glu Phe Asn Met Leu Ser Ala Leu Asn
 1905 1910 1915 1920
 Glu Ile Tyr Ser Tyr Val Ser Lys Tyr Ser Glu Glu Leu Ile Gly Ala
 1925 1930 1935
 Leu Glu Gln Asp Glu Gln Ala Arg Arg Gln Arg Leu Ala Tyr Lys Val
 1940 1945 1950
 Glu Gln Leu Ile Asn Ala Met Ser Ile Glu Ser
 1955 1960

<210> 44
 <211> 1905
 <212> PRT
 <213> *Xenopus laevis*

<400> 44
 Met Leu Leu His Ala Glu Arg Pro Leu Pro Phe His Leu Trp Thr Phe
 1 5 10 15
 Leu Val Leu Leu Gly Ser Trp Ile Ala Thr Gly Asp Gly Ser Pro Lys
 20 25 30
 Asp Phe Arg Thr Phe Thr Gly Ser Asp Trp Ser Leu Thr His Leu Val
 35 40 45
 Val His Asn Lys Thr Gly Glu Val Tyr Val Gly Ala Ile Asn Arg Ile
 50 55 60
 Tyr Lys Leu Ser Asn Asn Leu Thr Leu Leu Arg Thr His Val Thr Gly
 65 70 75 80
 Pro Val Glu Asp Asn Glu Lys Cys Tyr Pro Pro Pro Ser Val Gln Ser

85					90					95					
Cys	Pro	His	Gly	Leu	Ile	Thr	Thr	Asn	Asn	Val	Asn	Lys	Leu	Leu	Leu
			100					105					110		
Ile	Asp	Tyr	Ser	Asp	Asn	Arg	Leu	Ile	Ala	Cys	Gly	Ser	Ala	Ser	Gln
		115					120					125			
Gly	Ile	Cys	Gln	Phe	Leu	Arg	Leu	Asp	Asp	Leu	Phe	Lys	Leu	Gly	Glu
		130					135					140			
Pro	His	His	Arg	Lys	Glu	His	Tyr	Leu	Ser	Ser	Val	Asn	Glu	Ser	Gly
145							150					155			160
Thr	Met	Ser	Gly	Val	Ile	Ile	Glu	Val	Pro	Asn	Gly	Gln	Asn	Lys	Leu
				165					170					175	
Phe	Val	Gly	Thr	Pro	Ile	Asp	Gly	Lys	Ser	Glu	Tyr	Phe	Pro	Thr	Leu
			180					185					190		
Ser	Ser	Arg	Lys	Leu	Leu	Gly	Asn	Glu	Glu	Asn	Ala	Glu	Met	Phe	Gly
		195					200					205			
Phe	Val	Tyr	Gln	Asp	Glu	Phe	Val	Ser	Ser	Gln	Leu	Lys	Ile	Pro	Ser
		210					215					220			
Asp	Thr	Leu	Ser	Lys	Phe	Pro	Thr	Phe	Asp	Ile	Tyr	Tyr	Val	Tyr	Ser
225							230					235			240
Phe	Ser	Ser	Glu	Gln	Phe	Val	Tyr	Tyr	Leu	Thr	Leu	Gln	Leu	Asp	Thr
				245					250					255	
Gln	Leu	Thr	Ser	Pro	Asp	Ser	Thr	Gly	Glu	Gln	Phe	Phe	Thr	Ser	Lys
			260					265					270		
Ile	Val	Arg	Leu	Cys	Val	Asp	Asp	Pro	Lys	Phe	Tyr	Ser	Tyr	Val	Glu
		275					280					285			
Phe	Pro	Ile	Gly	Cys	Met	Lys	Asp	Gly	Val	Glu	Tyr	Arg	Leu	Ile	Gln
		290					295					300			
Asp	Ala	Tyr	Leu	Ser	Lys	Pro	Gly	Lys	Arg	Leu	Ala	Lys	Glu	Leu	Gly
305							310					315			320
Ile	Ser	Glu	Arg	Glu	Asp	Ile	Leu	Phe	Thr	Val	Phe	Ser	Gln	Gly	Gln
				325					330					335	
Lys	Asn	Arg	Ile	Lys	Pro	Pro	Lys	Glu	Ser	Val	Leu	Cys	Leu	Phe	Thr
			340					345					350		
Leu	Lys	Lys	Ile	Lys	Asp	Lys	Ile	Lys	Glu	Arg	Ile	Gln	Ser	Cys	Tyr
		355					360					365			
Arg	Gly	Asp	Gly	Lys	Leu	Ser	Leu	Pro	Trp	Leu	Leu	Asn	Lys	Glu	Leu
		370					375					380			
Gly	Cys	Ile	Asn	Ser	Pro	Leu	Gln	Ile	Asp	Asp	Asn	Phe	Cys	Gly	Gln

385		390		395		400
Asp Phe Asn Gln Pro Leu Gly Gly Thr Val Thr Ile Glu Gly Thr Pro						
	405		410		415	
Leu Phe Leu Asp Lys Glu Asp Gly Met Thr Ser Val Ala Ala Tyr Asp						
	420		425		430	
Tyr Arg Gly His Thr Val Val Phe Ala Gly Thr Arg Ser Gly Arg Val						
	435		440		445	
Lys Lys Ile Leu Val Asp Leu Ser Ala Ser Ser Ser His Leu Val Gln						
	450		455		460	
Gln Tyr Glu Asn Val Val Val His Glu Gly Asn Ala Ile Leu Arg Asp						
465		470		475		480
Leu Val Leu Ser Pro Asp Arg Gln Tyr Ile Tyr Ala Met Thr Glu Lys						
	485		490		495	
Gln Val Thr Arg Val Pro Val Glu Ser Cys Glu Gln Tyr Glu Ser Cys						
	500		505		510	
Asp Thr Cys Leu Gly Ser Arg Asp Pro His Cys Gly Trp Cys Val Leu						
	515		520		525	
His Asn Met Cys Ser Arg Lys Asp Lys Cys Glu Arg Ala Asp Glu Leu						
	530		535		540	
His Arg Phe Thr Ser Asp Gln Arg Gln Cys Val Gln Leu Thr Val His						
545		550		555		560
Pro Lys Asn Ile Ser Val Thr Val Ser Glu Val Pro Met Val Leu Gln						
	565		570		575	
Ala Trp Asn Val Pro Asp Leu Ser Ala Gly Val Asn Cys Ser Phe Glu						
	580		585		590	
Asp Phe Thr Glu Met Glu Gly Arg Ile Leu Asp Gly Lys Ile Tyr Cys						
	595		600		605	
Thr Ser Pro Ser Ala Lys Glu Val Ile Pro Ile Thr Arg Gly His Gly						
	610		615		620	
Asp Lys Arg Val Val Lys Leu Tyr Leu Lys Ser Lys Glu Thr Gly Lys						
625		630		635		640
Lys Phe Ala Ser Val Asp Phe Val Phe Tyr Asn Cys Ser Val His Gln						
	645		650		655	
Ser Cys Leu Ser Cys Val Asn Gly Ser Phe Pro Cys His Trp Cys Lys						
	660		665		670	
Tyr Arg His Val Cys Thr His Asn Ala Ala Asp Cys Ser Phe Gln Glu						
	675		680		685	
Gly Arg Val Asn Met Ser Glu Asp Cys Pro Gln Ile Leu Pro Ser Ser						

995	1000	1005
Pro Ser Thr Gly Lys Ala Glu Ile Gln Ile Leu Ile Asn Arg Ala Thr 1010 1015 1020		
Met Asn Asn Ser Glu Val His Tyr Asn Tyr Thr Glu Asp Pro Thr Val 1025 1030 1035 1040		
Gln Lys Ile Glu Pro Glu Trp Ser Ile Ala Ser Gly Gly Thr Pro Leu 1045 1050 1055		
Ile Val Thr Gly Met Asn Leu Ala Thr Ile Lys Glu Pro Lys Ile Arg 1060 1065 1070		
Ala Lys Tyr Gly Asp Val Glu Lys Glu Asn Asn Cys Thr Leu Tyr Asn 1075 1080 1085		
Asp Thr Thr Met Val Cys Leu Ala Pro Ser Val Asp Asn Pro Leu Arg 1090 1095 1100		
Ser Pro Pro Glu Asn Gly Asp Arg Pro Asp Glu Ile Gly Phe Ile Met 1105 1110 1115 1120		
Asp Asn Val His Ala Leu Leu Ile Val Asn Thr Thr Ser Phe Leu Tyr 1125 1130 1135		
Tyr Pro Asp Pro Val Phe Glu Pro Leu Thr Ala Ser Gly Asn Leu Glu 1140 1145 1150		
Leu Lys Pro Ser Ser Pro Leu Ile Ile Lys Gly Arg Asn Leu Ile Pro 1155 1160 1165		
Ala Ala Pro Gly Asn Phe Arg Leu Asn Tyr Thr Val Leu Ile Gly Asp 1170 1175 1180		
Thr Pro Cys Ala Leu Thr Val Ser Glu Thr Gln Leu Leu Cys Glu Ser 1185 1190 1195 1200		
Pro Asn Leu Thr Gly Gln His Lys Val Thr Ile Lys Ala Gly Gly Phe 1205 1210 1215		
Glu Tyr Ser Pro Gly Thr Leu Gln Ile Tyr Ser Asp Ser Leu Leu Thr 1220 1225 1230		
Leu Pro Ala Ile Ile Gly Ile Gly Gly Gly Gly Gly Leu Leu Leu Leu 1235 1240 1245		
Ile Ile Ile Ile Val Leu Ile Ala Tyr Lys Arg Lys Ser Arg Asp Ala 1250 1255 1260		
Asp Arg Thr Leu Lys Arg Leu Gln Leu Gln Met Asp Asn Leu Glu Ser 1265 1270 1275 1280		
Arg Val Ala Leu Glu Cys Lys Glu Ala Phe Ala Glu Leu Gln Thr Asp 1285 1290 1295		
Ile His Glu Leu Thr Asn Asp Leu Asp Gly Ala Gly Ile Pro Phe Leu		

1300	1305	1310
Glu Tyr Arg Thr Tyr Ala Met Arg Val Leu Phe Pro Gly Ile Glu Asp 1315 1320 1325		
His Pro Val Leu Lys Glu Met Glu Val Gln Ala Asn Val Glu Lys Ser 1330 1335 1340		
Leu Thr Leu Phe Gly Gln Leu Leu Thr Lys Lys His Phe Leu Leu Thr 1345 1350 1355 1360		
Phe Ile Arg Thr Leu Glu Ala Gln Arg Ser Phe Ser Met Arg Asp Arg 1365 1370 1375		
Gly Asn Val Ala Ser Leu Ile Met Thr Ala Leu Gln Gly Glu Met Glu 1380 1385 1390		
Tyr Ala Thr Gly Val Leu Lys Gln Leu Leu Ser Asp Leu Ile Glu Lys 1395 1400 1405		
Asn Leu Glu Ser Lys Asn His Pro Lys Leu Leu Leu Arg Arg Thr Glu 1410 1415 1420		
Ser Val Ala Glu Lys Met Leu Thr Asn Trp Phe Thr Phe Leu Leu Tyr 1425 1430 1435 1440		
Lys Phe Leu Lys Glu Cys Ala Gly Glu Pro Leu Phe Met Leu His Cys 1445 1450 1455		
Ala Ile Lys Gln Gln Met Glu Lys Gly Pro Ile Asp Ala Ile Thr Gly 1460 1465 1470		
Glu Ala Arg Tyr Ser Leu Ser Glu Asp Lys Leu Ile Arg Gln Gln Ile 1475 1480 1485		
Asp Tyr Lys Thr Leu Asn Pro Cys Ala Asp Asp Val Gly Leu Ser Asp 1490 1495 1500		
Glu Ser Cys Cys Arg Ser Pro Gln Thr Leu Asn Cys Val Asn Pro Glu 1505 1510 1515 1520		
Asn Glu Asn Ala Pro Glu Ile Pro Val Lys Val Leu Asn Cys Asp Thr 1525 1530 1535		
Ile Thr Gln Val Lys Glu Lys Leu Leu Asp Ala Val Tyr Lys Gly Val 1540 1545 1550		
Pro Tyr Ser Gln Arg Pro Lys Ala Gly Asp Met Asp Leu Glu Trp Arg 1555 1560 1565		
Gln Gly Arg Met Ala Arg Ile Ile Leu Gln Asp Glu Asp Val Thr Thr 1570 1575 1580		
Lys Ile Asp Asn Asp Trp Lys Arg Leu Asn Thr Leu Ala His Tyr Gln 1585 1590 1595 1600		
Val Thr Asp Gly Ser Ser Val Ala Leu Val Pro Lys Gln Asn Ser Ala		

1605					1610					1615						
Tyr	Asn	Ile	Ser	Asn	Ser	Ser	Thr	Phe	Thr	Lys	Ser	Leu	Ser	Arg	Tyr	
1620					1625					1630						
Glu	Ser	Met	Leu	Arg	Thr	Ala	Ser	Ser	Pro	Asp	Ser	Leu	Arg	Ser	Arg	
1635					1640					1645						
Thr	Pro	Met	Ile	Thr	Pro	Asp	Leu	Glu	Ser	Gly	Thr	Lys	Leu	Trp	His	
1650					1655					1660						
Leu	Val	Lys	Asn	His	Asp	His	Leu	Asp	Gln	Arg	Glu	Gly	Asp	Arg	Gly	
1665					1670					1675					1680	
Ser	Lys	Met	Val	Ser	Glu	Ile	Tyr	Leu	Thr	Arg	Leu	Leu	Ala	Thr	Lys	
1685					1690					1695						
Gly	Thr	Leu	Gln	Lys	Phe	Val	Asp	Asp	Leu	Phe	Glu	Thr	Ile	Phe	Ser	
1700					1705					1710						
Thr	Ala	His	Arg	Gly	Ser	Ala	Leu	Pro	Leu	Ala	Ile	Lys	Tyr	Met	Phe	
1715					1720					1725						
Asp	Phe	Leu	Asp	Glu	Gln	Ala	Asp	Lys	His	Gln	Ile	Thr	Asp	Tyr	Asp	
1730					1735					1740						
Val	Arg	His	Thr	Trp	Lys	Ser	Asn	Cys	Leu	Pro	Leu	Arg	Phe	Trp	Val	
1745					1750					1755					1760	
Asn	Val	Ile	Lys	Asn	Pro	Gln	Phe	Val	Phe	Asp	Ile	His	Lys	Asn	Ser	
1765					1770					1775						
Ile	Thr	Asp	Ala	Cys	Leu	Ser	Val	Val	Ala	Gln	Thr	Phe	Met	Asp	Ser	
1780					1785					1790						
Cys	Ser	Thr	Ser	Glu	His	Lys	Leu	Gly	Lys	Asp	Ser	Pro	Ser	Asn	Lys	
1795					1800					1805						
Leu	Leu	Tyr	Ala	Lys	Asp	Ile	Pro	Asn	Tyr	Lys	Ser	Trp	Val	Glu	Arg	
1810					1815					1820						
Tyr	Tyr	Ala	Asp	Ile	Ala	Lys	Met	Pro	Val	Ile	Ser	Asp	Gln	Asp	Met	
1825					1830					1835					1840	
Ser	Ala	Tyr	Leu	Ala	Glu	Gln	Ser	Arg	Leu	His	Leu	Ser	Gln	Phe	Asn	
1845					1850					1855						
Ser	Met	Ser	Ala	Leu	His	Glu	Ile	Tyr	Ser	Tyr	Ile	Thr	Lys	Tyr	Arg	
1860					1865					1870						
Asp	Glu	Ile	Leu	Thr	Ala	Leu	Glu	Lys	Asp	Glu	Gln	Ala	Arg	Arg	Gln	
1875					1880					1885						
Arg	Leu	Arg	Ser	Lys	Leu	Glu	Gln	Val	Ile	Asp	Thr	Met	Ala	Gln	Ser	
1890					1895					1900						

Ser

1905

<210> 45

<211> 813

<212> PRT

<213> Homo sapiens

<400> 45

Gly Thr Arg Val Val Cys Lys Val Val Asn Thr Thr Thr Leu Thr Cys
1 5 10 15

Leu Ala Pro Ser Leu Thr Thr Asp Tyr Arg Pro Gly Leu Asp Thr Val
20 25 30

Glu Arg Pro Asp Glu Phe Gly Phe Val Phe Asn Asn Val Gln Ser Leu
35 40 45

Leu Ile Tyr Asn Asp Thr Lys Phe Ile Tyr Tyr Pro Asn Pro Thr Phe
50 55 60

Glu Leu Leu Ser Pro Thr Gly Val Leu Asp Gln Lys Pro Gly Ser Pro
65 70 75 80

Ile Ile Leu Lys Gly Lys Asn Leu Cys Pro Pro Ala Ser Gly Gly Ala
85 90 95

Lys Leu Asn Tyr Thr Val Leu Ile Gly Glu Thr Pro Cys Ala Val Thr
100 105 110

Val Ser Glu Thr Gln Leu Leu Cys Glu Pro Pro Asn Leu Thr Gly Gln
115 120 125

His Lys Val Met Val His Val Gly Gly Met Val Phe Ser Pro Gly Ser
130 135 140

Val Ser Val Ile Ser Asp Ser Leu Leu Thr Leu Pro Ala Ile Val Ser
145 150 155 160

Ile Ala Ala Gly Gly Ser Leu Leu Leu Ile Ile Val Ile Ile Val Leu
165 170 175

Ile Ala Tyr Lys Arg Lys Ser Arg Glu Asn Asp Leu Thr Leu Lys Arg
180 185 190

Leu Gln Met Gln Met Asp Asn Leu Glu Ser Arg Val Ala Leu Glu Cys
195 200 205

Lys Glu Ala Phe Ala Glu Leu Gln Thr Asp Ile Asn Glu Leu Thr Ser
210 215 220

Asp Leu Asp Arg Ser Gly Ile Pro Tyr Leu Asp Tyr Arg Thr Tyr Ala
225 230 235 240

Met Arg Val Leu Phe Pro Gly Ile Glu Asp His Pro Val Leu Arg Glu
245 250 255

Leu	Glu	Val	Gln	Gly	Asn	Gly	Gln	Gln	His	Val	Glu	Lys	Ala	Leu	Lys	
			260					265					270			
Leu	Phe	Ala	Gln	Leu	Ile	Asn	Asn	Lys	Val	Phe	Leu	Leu	Thr	Phe	Ile	
		275					280					285				
Arg	Thr	Leu	Glu	Leu	Gln	Arg	Ser	Phe	Ser	Met	Arg	Asp	Arg	Gly	Asn	
	290					295					300					
Val	Ala	Ser	Leu	Ile	Met	Thr	Gly	Leu	Gln	Gly	Arg	Leu	Glu	Tyr	Ala	
305					310					315					320	
Thr	Asp	Val	Leu	Lys	Gln	Leu	Leu	Ser	Asp	Leu	Ile	Asp	Lys	Asn	Leu	
				325					330					335		
Glu	Asn	Lys	Asn	His	Pro	Lys	Leu	Leu	Leu	Arg	Arg	Thr	Glu	Ser	Val	
			340					345					350			
Ala	Glu	Lys	Met	Leu	Thr	Asn	Trp	Phe	Ala	Phe	Leu	Leu	His	Lys	Phe	
		355					360					365				
Leu	Lys	Glu	Cys	Ala	Gly	Glu	Pro	Leu	Phe	Met	Leu	Tyr	Cys	Ala	Ile	
	370					375					380					
Lys	Gln	Gln	Met	Glu	Lys	Gly	Pro	Ile	Asp	Ala	Ile	Thr	Gly	Glu	Ala	
385					390					395					400	
Arg	Tyr	Ser	Leu	Ser	Glu	Asp	Lys	Leu	Ile	Arg	Gln	Gln	Ile	Glu	Tyr	
			405						410					415		
Lys	Thr	Leu	Ile	Leu	Asn	Cys	Val	Asn	Pro	Asp	Asn	Glu	Asn	Ser	Pro	
			420					425					430			
Glu	Ile	Pro	Val	Lys	Val	Leu	Asn	Cys	Asp	Thr	Ile	Thr	Gln	Val	Lys	
	435						440					445				
Glu	Lys	Ile	Leu	Asp	Ala	Val	Tyr	Lys	Asn	Val	Pro	Tyr	Ser	Gln	Arg	
	450					455					460					
Pro	Arg	Ala	Val	Asp	Met	Asp	Leu	Glu	Trp	Arg	Gln	Gly	Arg	Ile	Ala	
465					470					475					480	
Arg	Val	Val	Leu	Gln	Asp	Glu	Asp	Ile	Thr	Thr	Lys	Ile	Glu	Gly	Asp	
			485						490					495		
Trp	Lys	Arg	Leu	Asn	Thr	Leu	Met	His	Tyr	Gln	Val	Ser	Asp	Arg	Ser	
			500					505					510			
Val	Val	Ala	Leu	Val	Pro	Lys	Gln	Thr	Ser	Ser	Tyr	Asn	Ile	Pro	Ala	
		515					520					525				
Ser	Ala	Ser	Ile	Ser	Arg	Thr	Ser	Ile	Ser	Arg	Tyr	Asp	Ser	Ser	Phe	
	530					535					540					
Arg	Tyr	Thr	Gly	Ser	Pro	Asp	Ser	Leu	Arg	Ser	Arg	Ala	Pro	Met	Ile	
545					550					555					560	

Thr Pro Asp Leu Glu Ser Gly Val Lys Val Trp His Leu Val Lys Asn
 565 570 575
 His Asp His Gly Asp Gln Lys Glu Gly Asp Arg Gly Ser Lys Met Val
 580 585 590
 Ser Glu Ile Tyr Leu Thr Arg Leu Leu Ala Thr Lys Gly Thr Leu Gln
 595 600 605
 Lys Phe Val Asp Asp Leu Phe Glu Thr Leu Leu Ser Thr Val His Arg
 610 615 620
 Gly Ser Ala Leu Pro Leu Ala Ile Lys Tyr Met Phe Asp Phe Leu Asp
 625 630 635 640
 Glu Gln Ala Asp Arg His Ser Ile His Asp Thr Asp Val Arg His Thr
 645 650 655
 Trp Lys Ser Asn Cys Leu Pro Leu Arg Phe Trp Val Asn Val Ile Lys
 660 665 670
 Asn Pro Gln Phe Val Phe Asp Ile His Lys Gly Ser Ile Thr Asp Ala
 675 680 685
 Cys Leu Ser Val Val Ala Gln Thr Phe Met Asp Ser Cys Ser Thr Ser
 690 695 700
 Glu His Arg Leu Gly Lys Asp Ser Pro Ser Asn Lys Leu Leu Tyr Ala
 705 710 715 720
 Lys Asp Ile Pro Ser Tyr Lys Ser Trp Val Glu Arg Tyr Tyr Ala Asp
 725 730 735
 Ile Ala Lys Leu Pro Ala Ile Ser Asp Gln Asp Met Asn Ala Tyr Leu
 740 745 750
 Ala Glu Gln Ser Arg Leu His Ala Val Glu Phe Asn Met Leu Ser Ala
 755 760 765
 Leu Asn Glu Ile Tyr Ser Tyr Val Ser Lys Tyr Ser Glu Glu Leu Ile
 770 775 780
 Gly Ala Leu Glu Gln Asp Glu Gln Ala Arg Arg Gln Arg Leu Ala Tyr
 785 790 795 800
 Lys Val Glu Gln Leu Ile Asn Ala Met Ser Ile Glu Ser
 805 810

<210> 46

<211> 593

<212> PRT

<213> Homo sapiens

<400> 46

Val Glu Leu Thr Val Val Trp Asn Gly His Phe Asn Ile Asp Asn Pro
 1 5 10 15

Ala Gln Asn Lys Val His Leu Tyr Lys Cys Gly Ala Met Arg Glu Ser
 20 25 30
 Cys Gly Leu Cys Leu Lys Ala Asp Pro Asp Phe Ala Cys Gly Trp Cys
 35 40 45
 Gln Gly Pro Gly Gln Cys Thr Leu Arg Gln His Cys Pro Ala Gln Glu
 50 55 60
 Ser Gln Trp Leu Glu Leu Ser Gly Ala Lys Ser Lys Cys Thr Asn Pro
 65 70 75 80
 Arg Ile Thr Glu Ile Ile Pro Val Thr Gly Pro Arg Glu Gly Gly Thr
 85 90 95
 Lys Val Thr Ile Arg Gly Glu Asn Leu Gly Leu Glu Phe Arg Asp Ile
 100 105 110
 Ala Ser His Val Lys Val Ala Gly Val Glu Cys Ser Pro Leu Val Asp
 115 120 125
 Gly Tyr Ile Pro Ala Glu Gln Ile Val Cys Glu Met Gly Glu Ala Lys
 130 135 140
 Pro Ser Gln His Ala Gly Phe Val Glu Ile Cys Val Ala Val Cys Arg
 145 150 155 160
 Pro Glu Phe Met Ala Arg Ser Ser Gln Leu Tyr Tyr Phe Met Thr Leu
 165 170 175
 Thr Leu Ser Asp Leu Lys Pro Ser Arg Gly Pro Met Ser Gly Gly Thr
 180 185 190
 Gln Val Thr Ile Thr Gly Thr Asn Leu Asn Ala Gly Ser Asn Val Val
 195 200 205
 Val Met Phe Gly Lys Gln Pro Cys Leu Phe His Arg Arg Ser Pro Ser
 210 215 220
 Tyr Ile Val Cys Asn Thr Thr Ser Ser Asp Glu Val Leu Glu Met Lys
 225 230 235 240
 Val Ser Val Gln Val Asp Arg Ala Lys Ile His Gln Asp Leu Val Phe
 245 250 255
 Gln Tyr Val Glu Asp Pro Thr Ile Val Arg Ile Glu Pro Glu Trp Ser
 260 265 270
 Ile Val Ser Gly Asn Thr Pro Ile Ala Val Trp Gly Thr His Leu Asp
 275 280 285
 Leu Ile Gln Asn Pro Gln Ile Arg Ala Lys His Gly Gly Lys Glu His
 290 295 300
 Ile Asn Ile Cys Glu Val Leu Asn Ala Thr Glu Met Thr Cys Gln Ala
 305 310 315 320

Pro Ala Leu Ala Leu Gly Pro Asp His Gln Ser Asp Leu Thr Glu Arg
 325 330 335
 Pro Glu Glu Phe Gly Phe Ile Leu Asp Asn Val Gln Ser Leu Leu Ile
 340 345 350
 Leu Asn Lys Thr Asn Phe Thr Tyr Tyr Pro Asn Pro Val Phe Glu Ala
 355 360 365
 Phe Gly Pro Ser Gly Ile Leu Glu Leu Lys Pro Gly Thr Pro Ile Ile
 370 375 380
 Leu Lys Gly Lys Asn Leu Ile Pro Pro Val Ala Gly Gly Asn Val Lys
 385 390 395 400
 Leu Asn Tyr Thr Val Leu Val Gly Glu Lys Pro Cys Thr Val Thr Val
 405 410 415
 Ser Asp Val Gln Leu Leu Cys Glu Ser Pro Asn Leu Ile Gly Arg His
 420 425 430
 Lys Val Met Ala Arg Val Gly Gly Met Glu Tyr Ser Pro Gly Met Val
 435 440 445
 Tyr Ile Ala Pro Asp Ser Pro Leu Ser Leu Pro Ala Ile Val Ser Ile
 450 455 460
 Ala Val Ala Gly Gly Leu Leu Ile Ile Phe Ile Val Ala Val Leu Ile
 465 470 475 480
 Ala Tyr Lys Arg Lys Ser Arg Glu Ser Asp Leu Thr Leu Lys Arg Leu
 485 490 495
 Gln Met Gln Met Asp Asn Leu Glu Ser Arg Val Ala Leu Glu Cys Lys
 500 505 510
 Glu Gly Thr Glu Trp Pro His Ala Gly Gly His Val Cys Val Arg Val
 515 520 525
 Cys Ile Cys Val Cys Met His Ile Cys Val Cys Val Cys Ile Cys Phe
 530 535 540
 Ile Tyr Lys Gln Ala Gly Trp Ala Ala Val Gly Ser Ala Gly Gly Trp
 545 550 555 560
 Arg Cys Val Cys Leu Cys Glu Cys Val Cys Val His Val Cys Val Cys
 565 570 575
 Thr Ser Val Cys Ile Tyr Val Ser Tyr Thr Ser Lys Gln Ala Gly Gln
 580 585 590
 Gln

<210> 47

<211> 477
 <212> PRT
 <213> Homo sapiens

<400> 47

Met	Leu	Pro	Pro	Gly	Ser	Asn	Gly	Thr	Ala	Tyr	Pro	Gly	Gln	Phe	Ala	1	5	10	15
Leu	Tyr	Gln	Gln	Leu	Ala	Gln	Gly	Asn	Ala	Val	Gly	Gly	Ser	Ala	Gly	20	25	30	
Ala	Pro	Pro	Leu	Gly	Pro	Ser	Gln	Val	Val	Thr	Ala	Cys	Leu	Leu	Thr	35	40	45	
Leu	Leu	Ile	Ile	Trp	Thr	Leu	Leu	Gly	Asn	Val	Leu	Val	Cys	Ala	Ala	50	55	60	
Ile	Val	Arg	Ser	Arg	His	Leu	Arg	Ala	Asn	Met	Thr	Asn	Val	Phe	Ile	65	70	75	80
Val	Ser	Leu	Ala	Val	Ser	Asp	Leu	Phe	Val	Ala	Leu	Leu	Val	Met	Pro	85	90	95	
Trp	Lys	Ala	Val	Ala	Glu	Val	Ala	Gly	Tyr	Trp	Pro	Phe	Gly	Ala	Phe	100	105	110	
Cys	Asp	Val	Trp	Val	Ala	Phe	Asp	Ile	Met	Cys	Ser	Thr	Ala	Ser	Ile	115	120	125	
Leu	Asn	Leu	Cys	Val	Ile	Ser	Val	Asp	Arg	Tyr	Trp	Ala	Ile	Ser	Arg	130	135	140	
Pro	Phe	Arg	Tyr	Lys	Arg	Lys	Met	Thr	Gln	Arg	Met	Ala	Leu	Val	Met	145	150	155	160
Val	Gly	Leu	Ala	Trp	Thr	Leu	Ser	Ile	Leu	Ile	Ser	Phe	Ile	Pro	Val	165	170	175	
Gln	Leu	Asn	Trp	His	Arg	Asp	Gln	Ala	Ala	Ser	Trp	Gly	Gly	Leu	Asp	180	185	190	
Leu	Pro	Asn	Asn	Leu	Ala	Asn	Trp	Thr	Pro	Trp	Glu	Glu	Asp	Phe	Trp	195	200	205	
Glu	Pro	Asp	Val	Asn	Ala	Glu	Asn	Cys	Asp	Ser	Ser	Leu	Asn	Arg	Thr	210	215	220	
Tyr	Ala	Ile	Ser	Ser	Ser	Leu	Ile	Ser	Phe	Tyr	Ile	Pro	Val	Ala	Ile	225	230	235	240
Met	Ile	Val	Thr	Tyr	Thr	Arg	Ile	Tyr	Arg	Ile	Ala	Gln	Val	Gln	Ile	245	250	255	
Arg	Arg	Ile	Ser	Ser	Leu	Glu	Arg	Ala	Ala	Glu	His	Ala	Gln	Ser	Cys	260	265	270	
Arg	Ser	Ser	Ala	Ala	Cys	Ala	Pro	Asp	Thr	Ser	Leu	Arg	Ala	Ser	Ile				

275	280	285
Lys Lys Glu Thr Lys Val	Leu Lys Thr Leu Ser Val	Ile Met Gly Val
290	295	300
Phe Val Cys Cys Trp Leu Pro Phe Phe Ile Leu Asn Cys Met Val Pro		
305	310	315 320
Phe Cys Ser Gly His Pro Glu Gly Pro Pro Ala Gly Phe Pro Cys Val		
	325	330 335
Ser Glu Thr Thr Phe Asp Val Phe Val Trp Phe Gly Trp Ala Asn Ser		
	340	345 350
Ser Leu Asn Pro Val Ile Tyr Ala Phe Asn Ala Asp Phe Gln Lys Val		
	355	360 365
Phe Ala Gln Leu Leu Gly Cys Ser His Phe Cys Ser Arg Thr Pro Val		
	370	375 380
Glu Thr Val Asn Ile Ser Asn Glu Leu Ile Ser Tyr Asn Gln Asp Ile		
385	390	395 400
Val Phe His Lys Glu Ile Ala Ala Ala Tyr Ile His Met Met Pro Asn		
	405	410 415
Ala Val Thr Pro Gly Asn Arg Glu Val Asp Asn Asp Glu Glu Glu Gly		
	420	425 430
Pro Phe Asp Arg Met Phe Gln Ile Tyr Gln Thr Ser Pro Asp Gly Asp		
	435	440 445
Pro Val Ala Glu Ser Val Trp Glu Leu Asp Cys Glu Gly Glu Ile Ser		
	450	455 460
Leu Asp Lys Ile Thr Pro Phe Thr Pro Asn Gly Phe His		
465	470	475

<210> 48

<211> 475

<212> PRT

<213> Homo sapiens

<400> 48

Met	Leu	Pro	Pro	Gly	Arg	Asn	Arg	Thr	Ala	Gln	Pro	Ala	Arg	Leu	Gly
1				5					10					15	

Leu	Gln	Arg	Gln	Leu	Ala	Gln	Val	Asp	Ala	Pro	Ala	Gly	Ser	Ala	Thr
			20					25					30		

Pro	Leu	Gly	Pro	Ala	Gln	Val	Val	Thr	Ala	Gly	Leu	Leu	Thr	Leu	Leu
		35						40					45		

Ile	Val	Trp	Thr	Leu	Leu	Gly	Asn	Val	Leu	Val	Cys	Ala	Ala	Ile	Val
				50			55				60				

Arg	Ser	Arg	His	Leu	Arg	Ala	Lys	Met	Thr	Asn	Ile	Phe	Ile	Val	Ser	
65					70					75					80	
Leu	Ala	Val	Ser	Asp	Leu	Phe	Val	Ala	Leu	Leu	Val	Met	Pro	Trp	Lys	
				85					90					95		
Ala	Val	Ala	Glu	Val	Ala	Gly	Tyr	Trp	Pro	Phe	Gly	Thr	Phe	Cys	Asp	
			100					105					110			
Ile	Trp	Val	Ala	Phe	Asp	Ile	Met	Cys	Ser	Thr	Ala	Ser	Ile	Leu	Asn	
		115					120					125				
Leu	Cys	Ile	Ile	Ser	Val	Asp	Arg	Tyr	Trp	Ala	Ile	Ser	Arg	Pro	Phe	
	130					135					140					
Arg	Tyr	Glu	Arg	Lys	Met	Thr	Gln	Arg	Val	Ala	Leu	Val	Met	Val	Gly	
145					150					155					160	
Leu	Ala	Trp	Thr	Leu	Ser	Ile	Leu	Ile	Ser	Phe	Ile	Pro	Val	Gln	Leu	
				165					170					175		
Asn	Trp	His	Arg	Asp	Lys	Ala	Gly	Ser	Gln	Gly	Gln	Glu	Gly	Leu	Leu	
			180					185					190			
Ser	Asn	Gly	Thr	Pro	Trp	Glu	Glu	Gly	Trp	Glu	Leu	Glu	Gly	Arg	Thr	
		195					200					205				
Glu	Asn	Cys	Asp	Ser	Ser	Leu	Asn	Arg	Thr	Tyr	Ala	Ile	Ser	Ser	Ser	
	210					215					220					
Leu	Ile	Ser	Phe	Tyr	Ile	Pro	Val	Ala	Ile	Met	Ile	Val	Thr	Tyr	Thr	
225					230					235					240	
Arg	Ile	Tyr	Arg	Ile	Ala	Gln	Val	Gln	Ile	Arg	Arg	Ile	Ser	Ser	Leu	
				245					250					255		
Glu	Arg	Ala	Ala	Glu	His	Ala	Gln	Ser	Cys	Arg	Ser	Arg	Gly	Ala	Tyr	
			260					265					270			
Glu	Pro	Asp	Pro	Ser	Leu	Arg	Ala	Ser	Ile	Lys	Lys	Glu	Thr	Lys	Val	
		275					280					285				
Phe	Lys	Thr	Leu	Ser	Met	Ile	Met	Gly	Val	Phe	Val	Cys	Cys	Trp	Leu	
	290					295					300					
Pro	Phe	Phe	Ile	Leu	Asn	Cys	Met	Val	Pro	Phe	Cys	Ser	Ser	Gly	Asp	
305					310					315					320	
Ala	Glu	Gly	Pro	Lys	Thr	Gly	Phe	Pro	Cys	Val	Ser	Glu	Thr	Thr	Phe	
				325					330					335		
Asp	Ile	Phe	Val	Trp	Phe	Gly	Trp	Ala	Asn	Ser	Ser	Leu	Asn	Pro	Ile	
			340					345					350			
Ile	Tyr	Ala	Phe	Asn	Ala	Asp	Phe	Arg	Lys	Val	Phe	Ala	Gln	Leu	Leu	
		355					360					365				

Gly Cys Ser His Phe Cys Phe Arg Thr Pro Val Gln Thr Val Asn Ile
370 375 380

Ser Asn Glu Leu Ile Ser Tyr Asn Gln Asp Thr Val Phe His Lys Glu
385 390 395 400

Ile Ala Thr Ala Tyr Val His Met Ile Pro Asn Ala Val Ser Ser Gly
405 410 415

Asp Arg Glu Val Gly Glu Glu Glu Glu Gly Pro Phe Asp His Met
420 425 430

Ser Gln Ile Ser Pro Thr Thr Pro Asp Gly Asp Leu Ala Ala Glu Ser
435 440 445

Val Trp Glu Leu Asp Cys Glu Glu Glu Val Ser Leu Gly Lys Ile Ser
450 455 460

Pro Leu Thr Pro Asn Cys Phe Asp Lys Thr Ala
465 470 475

<210> 49
<211> 457
<212> PRT
<213> Homo sapiens

<400> 49

Met Tyr Gln Pro Phe Gln His Leu Asp Ser Asp Gln Val Ala Ser Trp
1 5 10 15

Gln Ser Pro Glu Met Leu Met Asn Lys Ser Val Ser Arg Glu Ser Gln
20 25 30

Arg Arg Lys Glu Leu Val Ala Gly Gln Ile Val Thr Gly Ser Leu Leu
35 40 45

Leu Leu Leu Ile Phe Trp Thr Leu Phe Gly Asn Ile Leu Val Cys Thr
50 55 60

Ala Val Met Arg Phe Arg His Leu Arg Ser Arg Val Thr Asn Ile Phe
65 70 75 80

Ile Val Ser Leu Ala Val Ser Asp Leu Leu Val Ala Leu Leu Val Met
85 90 95

Pro Trp Lys Ala Val Ala Glu Val Ala Gly His Trp Pro Phe Gly Ala
100 105 110

Phe Cys Asp Ile Trp Val Ala Phe Asp Ile Met Cys Ser Thr Ala Ser
115 120 125

Ile Leu Asn Leu Cys Val Ile Ser Val Asp Arg Tyr Trp Ala Ile Ser
130 135 140

Ser Pro Phe Arg Tyr Glu Arg Lys Met Thr Gln Arg Val Ala Leu Leu
145 150 155 160

Met Ile Ser Thr Ala Trp Ala Leu Ser Val Leu Ile Ser Phe Ile Pro
 165 170 175
 Val Gln Leu Ser Trp His Lys Ser Glu Thr Glu Asp His Leu Leu Ser
 180 185 190
 Asn His Ser Thr Gly Asn Cys Asp Ser Ser Leu Asn Arg Thr Tyr Ala
 195 200 205
 Ile Ser Ser Ser Leu Ile Ser Phe Tyr Ile Pro Val Ala Ile Met Ile
 210 215 220
 Val Thr Tyr Thr Arg Ile Tyr Arg Ile Ala Gln Ile Gln Ile Lys Arg
 225 230 235 240
 Ile Ser Thr Leu Glu Arg Ala Ala Glu His Ala Gln Ser Cys Arg Ser
 245 250 255
 Asn Arg Val Asp Ser Cys Ser Arg His His Gln Thr Ser Leu Arg Thr
 260 265 270
 Ser Ile Lys Lys Glu Thr Lys Val Leu Lys Thr Leu Ser Ile Ile Met
 275 280 285
 Gly Val Phe Val Cys Cys Trp Leu Pro Phe Phe Ile Leu Asn Cys Met
 290 295 300
 Val Pro Phe Cys Asp Arg Ser Pro Gly His Pro Gln Ala Gly Leu Pro
 305 310 315 320
 Cys Val Ser Glu Thr Thr Phe Asp Ile Phe Val Trp Phe Gly Trp Ala
 325 330 335
 Asn Ser Ser Leu Asn Pro Ile Ile Tyr Ala Phe Asn Ala Asp Phe Arg
 340 345 350
 Lys Val Phe Ser Ser Leu Leu Gly Cys Gly His Trp Cys Ser Thr Thr
 355 360 365
 Pro Val Glu Thr Val Asn Ile Ser Asn Glu Leu Ile Ser Tyr Asn Gln
 370 375 380
 Asp Thr Leu Phe His Lys Asp Ile Val Thr Ala Tyr Val Asn Met Ile
 385 390 395 400
 Pro Asn Val Val Asp Cys Ile Asp Asp Asn Glu Asp Ala Phe Asp His
 405 410 415
 Met Ser Gln Ile Ser Gln Thr Ser Ala Asn Asn Glu Leu Ala Thr Asp
 420 425 430
 Ser Met Cys Glu Leu Asp Ser Glu Val Asp Ile Ser Leu His Lys Ile
 435 440 445
 Thr Pro Ser Met Ser Asn Gly Ile His
 450 455

<210> 50
 <211> 486
 <212> PRT
 <213> Gallus gallus

<400> 50

Met	Leu	Arg	Gly	Gly	Arg	Ser	Pro	Leu	Pro	Pro	Pro	Ala	Gly	Pro	Pro
1				5					10					15	
Gly	Gly	Ala	Arg	Gly	Gln	Ala	Gly	Ala	Gly	Ala	Ala	Gln	Val	Ala	Ala
			20					25					30		
Gly	Ser	Leu	Leu	Ala	Leu	Leu	Ile	Leu	Trp	Thr	Leu	Phe	Gly	Asn	Val
		35					40					45			
Leu	Val	Cys	Ala	Ala	Ile	Val	Arg	Tyr	Arg	His	Leu	Arg	Ser	Lys	Val
	50					55					60				
Thr	Asn	Ile	Phe	Ile	Val	Ser	Leu	Ala	Val	Ser	Asp	Leu	Leu	Val	Ala
65					70					75					80
Val	Leu	Val	Met	Pro	Trp	Lys	Ala	Val	Ala	Glu	Val	Ala	Gly	Tyr	Trp
			85						90					95	
Pro	Phe	Gly	Ala	Phe	Gln	Asn	Val	Trp	Val	Ala	Phe	Asp	Ile	Met	Cys
			100					105					110		
Ser	Thr	Ala	Ser	Ile	Leu	Asn	Leu	Cys	Val	Ile	Ser	Val	Asp	Arg	Tyr
		115					120					125			
Trp	Ala	Ile	Ser	Ser	Pro	Phe	Arg	Tyr	Glu	Arg	Lys	Met	Thr	Gln	Arg
	130					135					140				
Leu	Ala	Leu	Val	Met	Ile	Gly	Val	Ala	Trp	Ala	Leu	Ser	Val	Leu	Ile
145					150				155						160
Ser	Phe	Ile	Pro	Val	Gln	Leu	Asn	Trp	His	Arg	Gly	Gly	Asp	Ala	Ala
			165					170					175		
Thr	Ala	Ala	Ala	Ala	Gly	Asp	Ile	Glu	Asp	Gly	Phe	Asp	Thr	Gly	Trp
			180					185					190		
Glu	Ala	Ala	Gly	Ala	Phe	Thr	Thr	Trp	Ala	Glu	Asp	Met	Ser	Thr	Thr
	195						200					205			
Trp	Val	Ala	Leu	Ala	Ala	Met	Thr	Pro	Ser	Glu	Gly	Thr	Ser	Gly	Ser
	210					215					220				
Asn	Asn	Thr	Val	Pro	Gly	Pro	Ser	Glu	Ser	Cys	Asp	Ser	Ser	Leu	Asn
225					230					235					240
Arg	Thr	Tyr	Ala	Ile	Ser	Ser	Ser	Leu	Ile	Ser	Phe	Tyr	Ile	Pro	Val
			245					250					255		
Ala	Ile	Met	Ile	Val	Thr	Tyr	Thr	Arg	Ile	Tyr	Arg	Ile	Ala	Gln	Val

260					265					270					
Gln	Ile	Arg	Arg	Ile	Ser	Ser	Leu	Glu	Arg	Ala	Ala	Glu	His	Ala	Gln
	275						280					285			
Ser	Cys	Arg	Cys	Asn	His	Val	Asp	Cys	His	His	His	Thr	Ser	Leu	Lys
	290					295					300				
Ser	Ser	Ile	Arg	Lys	Glu	Thr	Lys	Val	Leu	Lys	Thr	Leu	Ser	Ile	Ile
305					310					315					320
Met	Gly	Val	Phe	Val	Cys	Cys	Trp	Leu	Pro	Phe	Phe	Ile	Leu	Asn	Cys
				325					330					335	
Met	Val	Pro	Phe	Cys	Glu	Ser	Pro	Pro	Ser	Asp	Pro	Arg	Ala	Gly	Leu
			340					345					350		
Pro	Cys	Val	Ser	Glu	Thr	Thr	Phe	Asn	Ile	Phe	Val	Trp	Phe	Gly	Trp
	355						360					365			
Ala	Asn	Ser	Ser	Leu	Asn	Pro	Ile	Ile	Tyr	Ala	Phe	Asn	Ala	Asp	Phe
	370					375					380				
Arg	Lys	Val	Phe	Ser	Asn	Leu	Leu	Gly	Cys	Gly	Gln	Phe	Cys	Ser	Ser
385					390					395					400
Thr	Pro	Val	Glu	Thr	Val	Asn	Ile	Ser	Asn	Glu	Leu	Ile	Ser	Tyr	His
			405					410					415		
Gln	Asp	Thr	Phe	His	Lys	Glu	Ile	Val	Thr	Ala	Tyr	Val	Asn	Met	Ile
		420						425					430		
Pro	Asn	Val	Val	Asp	Cys	Glu	Glu	Asn	Arg	Glu	Asp	Pro	Phe	Asp	Arg
	435						440					445			
Met	Ser	Gln	Ile	Ser	Pro	Asp	Pro	Glu	Val	Ala	Thr	Asp	Ser	Val	Cys
	450					455					460				
Glu	Leu	Asp	Cys	Glu	Gly	Glu	Ile	Ser	Leu	Gly	Lys	Ile	Thr	Pro	Phe
465					470					475					480
Thr	Pro	Asn	Gly	Leu	His										
				485											

<210> 51
 <211> 458
 <212> PRT
 <213> Anguilla anguilla

<400> 51
 Met Gly Ser Pro Ala Lys Tyr Leu Ser Val His Glu Thr Gln Ser Val
 1 5 10 15
 Pro Phe Phe Ile Gly Glu Ile Met Trp Asn Thr Ser Glu Ser Ala Glu
 20 25 30

Lys Thr Asp Gly Lys Lys Glu Leu Ile Val Arg Thr Val Thr Gly Cys
 35 40 45
 Leu Leu Ser Leu Leu Ile Leu Trp Thr Leu Leu Gly Asn Ile Leu Val
 50 55 60
 Cys Ser Ala Val Leu Lys Phe Arg His Leu Arg Thr Lys Val Thr Asn
 65 70 75 80
 Ile Phe Ile Val Ser Leu Ala Val Ser Asp Leu Phe Val Ala Val Leu
 85 90 95
 Val Met Pro Trp Lys Ala Val Ala Glu Val Ala Gly Tyr Trp Pro Phe
 100 105 110
 Gly Pro Phe Cys Asn Ile Trp Val Ala Phe Asp Ile Met Cys Ser Thr
 115 120 125
 Ala Ser Ile Leu Asn Leu Cys Ile Ile Ser Val Asp Arg Tyr Trp Ala
 130 135 140
 Ile Ser Ser Pro Phe Arg Tyr Glu Arg Lys Met Thr Gln Arg Val Ala
 145 150 155 160
 Phe Val Met Ile Ser Val Thr Trp Thr Leu Ser Val Leu Ile Ser Phe
 165 170 175
 Ile Pro Val Gln Leu Asn Trp His Lys Ala Ser Asp Glu Glu Val Trp
 180 185 190
 Ile Asn Gly Thr Ser Phe Gly Glu Lys Ser Glu Asn Cys Asp Ser Ser
 195 200 205
 Leu Asn Arg Glu Tyr Ala Ile Ser Ser Ser Leu Ile Ser Phe Tyr Ile
 210 215 220
 Pro Val Ala Ile Met Ile Val Thr Tyr Thr Arg Ile Tyr Arg Ile Ala
 225 230 235 240
 Gln Ile Gln Ile Arg Arg Ile Ser Ser Leu Glu Arg Ala Ala Glu His
 245 250 255
 Ala Gln Ser Cys Arg Thr Asn Arg Leu Glu Cys Gln His His Asn Thr
 260 265 270
 Leu Lys Thr Ser Ile Lys Arg Glu Thr Lys Val Phe Lys Thr Leu Ser
 275 280 285
 Val Ile Met Gly Val Phe Val Cys Cys Trp Leu Pro Phe Phe Ile Leu
 290 295 300
 Asn Cys Ile Val Pro Phe Cys Asp Arg Pro Pro Thr Asp His Thr Ala
 305 310 315 320
 Gly Leu Pro Cys Val Ser Asp Thr Thr Phe Asp Val Phe Val Trp Phe
 325 330 335

Gly Trp Thr Asn Ser Ser Leu Asn Pro Ile Ile Tyr Ala Phe Asn Ala
 340 345 350
 Asp Phe Arg Lys Ala Phe Ala Ser Leu Leu Gly Cys Arg Asn Phe Cys
 355 360 365
 Ser Arg Thr Pro Val Glu Thr Val Asn Ile Ser Asn Glu Leu Val Ser
 370 375 380
 Tyr Asn Gln Asp Thr Leu Phe His Lys Glu Ile Val Thr Ala Tyr Val
 385 390 395 400
 Asn Met Ile Pro Asn Val Val Asp Cys Ile Asp Asp Asn Glu Asp Thr
 405 410 415
 Phe Asp Arg Ile Ser Gln Phe Ser His Asn Asn Glu Ile Ala Thr Asp
 420 425 430
 Ser Val Cys Asp Leu Asp Asp Cys Glu Ala Asp Ile Cys Leu Asp Arg
 435 440 445
 Leu Ala Pro Phe Thr Pro Asn Gly Leu His
 450 455

<210> 52
 <211> 879
 <212> PRT
 <213> Rattus norvegicus

<400> 52
 Met Lys Met Leu Thr Arg Leu Gln Ile Leu Met Leu Ala Leu Phe Ser
 1 5 10 15
 Lys Gly Phe Leu Leu Ser Leu Gly Asp His Asn Phe Met Arg Arg Glu
 20 25 30
 Ile Lys Ile Glu Gly Asp Leu Val Leu Gly Gly Leu Phe Pro Ile Asn
 35 40 45
 Glu Lys Gly Thr Gly Thr Glu Glu Cys Gly Arg Ile Asn Glu Asp Arg
 50 55 60
 Gly Ile Gln Arg Leu Glu Ala Met Leu Phe Ala Ile Asp Glu Ile Asn
 65 70 75 80
 Lys Asp Asn Tyr Leu Leu Pro Gly Val Lys Leu Gly Val His Ile Leu
 85 90 95
 Asp Thr Cys Ser Arg Asp Thr Tyr Ala Leu Glu Gln Ser Leu Glu Phe
 100 105 110
 Val Arg Ala Ser Leu Thr Lys Val Asp Glu Ala Glu Tyr Met Cys Pro
 115 120 125
 Asp Gly Ser Tyr Ala Ile Gln Glu Asn Ile Pro Leu Leu Ile Ala Gly
 130 135 140

Val	Ile	Gly	Gly	Ser	Tyr	Ser	Ser	Val	Ser	Ile	Gln	Val	Ala	Asn	Leu	
145					150					155					160	
Leu	Arg	Leu	Phe	Gln	Ile	Pro	Gln	Ile	Ser	Tyr	Ala	Ser	Thr	Ser	Ala	
				165					170					175		
Lys	Leu	Ser	Asp	Lys	Ser	Arg	Tyr	Asp	Tyr	Phe	Ala	Arg	Thr	Val	Pro	
			180					185					190			
Pro	Asp	Phe	Tyr	Gln	Ala	Lys	Ala	Met	Ala	Glu	Ile	Leu	Arg	Phe	Phe	
		195					200					205				
Asn	Trp	Thr	Tyr	Val	Ser	Thr	Val	Ala	Ser	Glu	Gly	Asp	Tyr	Gly	Glu	
	210					215					220					
Thr	Gly	Ile	Glu	Ala	Phe	Glu	Gln	Glu	Ala	Arg	Leu	Arg	Asn	Ile	Cys	
225					230					235					240	
Ile	Ala	Thr	Ala	Glu	Lys	Val	Gly	Arg	Ser	Asn	Ile	Arg	Lys	Ser	Tyr	
				245					250					255		
Asp	Ser	Val	Ile	Arg	Glu	Leu	Leu	Gln	Lys	Pro	Asn	Ala	Arg	Val	Val	
			260					265					270			
Val	Leu	Phe	Met	Arg	Ser	Asp	Asp	Ser	Arg	Glu	Leu	Ile	Ala	Ala	Ala	
		275					280					285				
Asn	Arg	Val	Asn	Ala	Ser	Phe	Thr	Trp	Val	Ala	Ser	Asp	Gly	Trp	Gly	
	290					295					300					
Ala	Gln	Glu	Ser	Ile	Val	Lys	Gly	Ser	Glu	His	Val	Ala	Tyr	Gly	Ala	
305					310					315					320	
Ile	Thr	Leu	Glu	Leu	Ala	Ser	His	Pro	Val	Arg	Gln	Phe	Asp	Arg	Tyr	
				325					330					335		
Phe	Gln	Ser	Leu	Asn	Pro	Tyr	Asn	Asn	His	Arg	Asn	Pro	Trp	Phe	Arg	
			340					345					350			
Asp	Phe	Trp	Glu	Gln	Lys	Phe	Gln	Cys	Ser	Leu	Gln	Asn	Lys	Arg	Asn	
		355					360					365				
His	Arg	Gln	Val	Cys	Asp	Lys	His	Leu	Ala	Ile	Asp	Ser	Ser	Asn	Tyr	
					370		375				380					
Glu	Gln	Glu	Ser	Lys	Ile	Met	Phe	Val	Val	Asn	Ala	Val	Tyr	Ala	Met	
385					390					395					400	
Ala	His	Ala	Leu	His	Lys	Met	Gln	Arg	Thr	Leu	Cys	Pro	Asn	Thr	Thr	
				405					410					415		
Lys	Leu	Cys	Asp	Ala	Met	Lys	Ile	Leu	Asp	Gly	Lys	Lys	Leu	Tyr	Lys	
			420					425					430			
Glu	Tyr	Leu	Leu	Lys	Ile	Asn	Phe	Thr	Ala	Pro	Phe	Asn	Pro	Asn	Lys	
		435					440					445				

Gly	Ala	Asp	Ser	Ile	Val	Lys	Phe	Asp	Thr	Phe	Gly	Asp	Gly	Met	Gly	
450						455					460					
Arg	Tyr	Asn	Val	Phe	Asn	Leu	Gln	Gln	Thr	Gly	Gly	Lys	Tyr	Ser	Tyr	
465					470					475					480	
Leu	Lys	Val	Gly	His	Trp	Ala	Glu	Thr	Leu	Ser	Leu	Asp	Val	Asp	Ser	
				485					490					495		
Ile	His	Trp	Ser	Arg	Asn	Ser	Val	Pro	Thr	Ser	Gln	Cys	Ser	Asp	Pro	
			500					505					510			
Cys	Ala	Pro	Asn	Glu	Met	Lys	Asn	Met	Gln	Pro	Gly	Asp	Val	Cys	Cys	
		515					520					525				
Trp	Ile	Cys	Ile	Pro	Cys	Glu	Pro	Tyr	Glu	Tyr	Leu	Val	Asp	Glu	Phe	
530						535					540					
Thr	Cys	Met	Asp	Cys	Gly	Pro	Gly	Gln	Trp	Pro	Thr	Ala	Asp	Leu	Ser	
545					550					555					560	
Gly	Cys	Tyr	Asn	Leu	Pro	Glu	Asp	Tyr	Ile	Lys	Trp	Glu	Asp	Ala	Trp	
			565						570					575		
Ala	Ile	Gly	Pro	Val	Thr	Ile	Ala	Cys	Leu	Gly	Phe	Leu	Cys	Thr	Cys	
			580					585					590			
Ile	Val	Ile	Thr	Val	Phe	Ile	Lys	His	Asn	Asn	Thr	Pro	Leu	Val	Lys	
		595					600					605				
Ala	Ser	Gly	Arg	Glu	Leu	Cys	Tyr	Ile	Leu	Leu	Phe	Gly	Val	Ser	Leu	
610						615					620					
Ser	Tyr	Cys	Met	Thr	Phe	Phe	Phe	Ile	Ala	Lys	Pro	Ser	Pro	Val	Ile	
625					630					635					640	
Cys	Ala	Leu	Arg	Arg	Leu	Gly	Leu	Gly	Thr	Ser	Phe	Ala	Ile	Cys	Tyr	
				645					650					655		
Ser	Ala	Leu	Leu	Thr	Lys	Thr	Asn	Cys	Ile	Ala	Arg	Ile	Phe	Asp	Gly	
			660					665					670			
Val	Lys	Asn	Gly	Ala	Gln	Arg	Pro	Lys	Phe	Ile	Ser	Pro	Ser	Ser	Gln	
		675					680					685				
Val	Phe	Ile	Cys	Leu	Gly	Leu	Ile	Leu	Val	Gln	Ile	Val	Met	Val	Ser	
690						695					700					
Val	Trp	Leu	Ile	Leu	Glu	Thr	Pro	Gly	Thr	Arg	Arg	Tyr	Thr	Leu	Pro	
705					710					715					720	
Glu	Lys	Arg	Glu	Thr	Val	Ile	Leu	Lys	Cys	Asn	Val	Lys	Asp	Ser	Ser	
				725					730					735		
Met	Leu	Ile	Ser	Leu	Thr	Tyr	Asp	Val	Val	Leu	Val	Ile	Leu	Cys	Thr	
			740					745					750			

Val Tyr Ala Phe Lys Thr Arg Lys Cys Pro Glu Asn Phe Asn Glu Ala
 755 760 765
 Lys Phe Ile Gly Phe Thr Met Tyr Thr Thr Cys Ile Ile Trp Leu Ala
 770 775 780
 Phe Leu Pro Ile Phe Tyr Val Thr Ser Ser Asp Tyr Arg Val Gln Thr
 785 790 795 800
 Thr Thr Met Cys Ile Ser Val Ser Leu Ser Gly Phe Val Val Leu Gly
 805 810 815
 Cys Leu Phe Ala Pro Lys Val His Ile Val Leu Phe Gln Pro Gln Lys
 820 825 830
 Asn Val Val Thr His Arg Leu His Leu Asn Arg Phe Ser Val Ser Gly
 835 840 845
 Thr Ala Thr Thr Tyr Ser Gln Ser Ser Ala Ser Thr Tyr Val Pro Thr
 850 855 860
 Val Cys Asn Gly Arg Glu Val Leu Asp Ser Thr Thr Ser Ser Leu
 865 870 875

<210> 53
 <211> 879
 <212> PRT
 <213> Mus musculus

<400> 53
 Met Lys Met Leu Thr Arg Leu Gln Val Leu Met Leu Ala Leu Phe Ser
 1 5 10 15
 Lys Gly Phe Leu Val Ser Leu Gly Asp His Asn Phe Met Arg Arg Glu
 20 25 30
 Ile Lys Ile Glu Gly Asp Leu Val Leu Gly Gly Leu Phe Pro Ile Asn
 35 40 45
 Glu Lys Gly Thr Gly Thr Glu Glu Cys Gly Arg Ile Asn Glu Asp Arg
 50 55 60
 Gly Ile Gln Arg Leu Glu Ala Met Leu Phe Ala Ile Asp Glu Ile Asn
 65 70 75 80
 Lys Asp Asn Tyr Leu Leu Pro Gly Val Lys Leu Gly Val His Ile Leu
 85 90 95
 Asp Thr Cys Ser Arg Asp Thr Tyr Ala Leu Glu Gln Ser Leu Glu Phe
 100 105 110
 Val Arg Ala Ser Leu Thr Lys Val Asp Glu Ala Glu Tyr Met Cys Pro
 115 120 125
 Asp Gly Ser Tyr Ala Ile Gln Glu Asn Ile Pro Leu Leu Ile Ala Gly

130	135	140
Val Ile Gly Gly Ser Tyr Ser Ser Val Ser Ile Gln Val Ala Asn Leu 145 150 155 160		
Leu Arg Leu Phe Gln Ile Pro Gln Ile Ser Tyr Ala Ser Thr Ser Ala 165 170 175		
Lys Leu Ser Asp Lys Ser Arg Tyr Asp Tyr Phe Ala Arg Thr Val Pro 180 185 190		
Pro Asp Phe Tyr Gln Ala Lys Ala Met Ala Glu Ile Leu Arg Tyr Phe 195 200 205		
Asn Trp Thr Tyr Val Ser Thr Val Ala Ser Glu Gly Asp Tyr Gly Glu 210 215 220		
Thr Gly Ile Glu Ala Phe Glu Gln Glu Ala Arg Leu Arg Asn Ile Cys 225 230 235 240		
Ile Ala Thr Ala Glu Lys Val Gly Arg Ser Asn Ile Arg Lys Ser Tyr 245 250 255		
Asp Ser Val Ile Arg Glu Leu Leu Gln Lys Pro Asn Ala Arg Val Val 260 265 270		
Val Leu Phe Met Arg Ser Asp Asp Ser Arg Glu Leu Ile Ala Ala Ala 275 280 285		
Ser Arg Val Asn Ala Ser Phe Thr Trp Val Ala Ser Asp Gly Trp Gly 290 295 300		
Ala Gln Glu Ser Ile Val Lys Gly Ser Glu His Val Ala Tyr Gly Ala 305 310 315 320		
Ile Thr Leu Glu Leu Ala Ser His Pro Val Arg Gln Phe Asp Arg Tyr 325 330 335		
Phe Gln Ser Leu Asn Pro Tyr Asn Asn His Arg Asn Pro Trp Phe Arg 340 345 350		
Asp Phe Trp Glu Gln Lys Phe Gln Cys Ser Leu Gln Asn Lys Arg Asn 355 360 365		
His Arg Gln Ile Cys Asp Lys His Leu Ala Ile Asp Ser Ser Asn Tyr 370 375 380		
Glu Gln Glu Ser Lys Ile Met Phe Val Val Asn Ala Val Tyr Ala Met 385 390 395 400		
Ala His Ala Leu His Lys Met Gln Arg Thr Leu Cys Pro Asn Thr Thr 405 410 415		
Lys Leu Cys Asp Ala Met Lys Ile Leu Asp Gly Lys Lys Leu Tyr Lys 420 425 430		
Asp Tyr Leu Leu Lys Ile Asn Phe Thr Ala Pro Phe Asn Pro Asn Lys		

435					440					445					
Gly	Ala	Asp	Ser	Ile	Val	Lys	Phe	Asp	Thr	Tyr	Gly	Asp	Gly	Met	Gly
450						455					460				
Arg	Tyr	Asn	Val	Phe	Asn	Phe	Gln	His	Ile	Gly	Gly	Lys	Tyr	Ser	Tyr
465					470					475					480
Leu	Lys	Val	Gly	His	Trp	Ala	Glu	Thr	Leu	Tyr	Leu	Asp	Val	Asp	Ser
				485					490					495	
Ile	His	Trp	Ser	Arg	Asn	Ser	Val	Pro	Thr	Ser	Gln	Cys	Ser	Asp	Pro
			500					505					510		
Cys	Ala	Pro	Asn	Glu	Met	Lys	Asn	Met	Gln	Pro	Gly	Asp	Val	Cys	Cys
			515				520					525			
Trp	Ile	Cys	Ile	Pro	Cys	Glu	Pro	Tyr	Glu	Tyr	Leu	Val	Asp	Glu	Phe
	530					535					540				
Thr	Cys	Met	Asp	Cys	Gly	Pro	Gly	Gln	Trp	Pro	Thr	Ala	Asp	Leu	Ser
545					550					555					560
Gly	Cys	Tyr	Asn	Leu	Pro	Glu	Asp	Tyr	Ile	Arg	Trp	Glu	Asp	Ala	Trp
			565						570					575	
Ala	Ile	Gly	Pro	Val	Thr	Ile	Ala	Cys	Leu	Gly	Phe	Met	Cys	Thr	Cys
			580					585					590		
Ile	Val	Ile	Thr	Val	Phe	Ile	Lys	His	Asn	Asn	Thr	Pro	Leu	Val	Lys
		595					600					605			
Ala	Ser	Gly	Arg	Glu	Leu	Cys	Tyr	Ile	Leu	Leu	Phe	Gly	Val	Ser	Leu
	610					615					620				
Ser	Tyr	Cys	Met	Thr	Phe	Phe	Phe	Ile	Ala	Lys	Pro	Ser	Pro	Val	Ile
625					630					635					640
Cys	Ala	Leu	Arg	Arg	Leu	Gly	Leu	Gly	Thr	Ser	Phe	Ala	Ile	Cys	Tyr
			645						650					655	
Ser	Ala	Leu	Leu	Thr	Lys	Thr	Asn	Cys	Ile	Ala	Arg	Ile	Phe	Asp	Gly
			660					665					670		
Val	Lys	Asn	Gly	Ala	Gln	Arg	Pro	Lys	Phe	Ile	Ser	Pro	Ser	Ser	Gln
		675					680					685			
Val	Phe	Ile	Cys	Leu	Gly	Leu	Ile	Leu	Val	Gln	Ile	Val	Met	Val	Ser
	690					695					700				
Val	Trp	Leu	Ile	Leu	Glu	Thr	Pro	Gly	Thr	Arg	Arg	Tyr	Thr	Leu	Pro
705					710					715					720
Glu	Lys	Arg	Glu	Thr	Val	Ile	Leu	Lys	Cys	Asn	Val	Lys	Asp	Ser	Ser
			725						730					735	
Met	Leu	Ile	Ser	Leu	Thr	Tyr	Asp	Val	Val	Leu	Val	Ile	Leu	Cys	Thr

740	745	750
Val Tyr Ala Phe Lys Thr Arg Lys Cys Pro Glu Asn Phe Asn Glu Ala		
755	760	765
Lys Phe Ile Gly Phe Thr Met Tyr Thr Thr Cys Ile Ile Trp Leu Ala		
770	775	780
Phe Leu Pro Ile Phe Tyr Val Thr Ser Ser Asp Tyr Arg Val Gln Thr		
785	790	795
Thr Thr Met Cys Ile Ser Val Ser Leu Ser Gly Phe Val Val Leu Gly		
	805	810
		815
Cys Leu Phe Ala Pro Lys Val His Ile Val Leu Phe Gln Pro Gln Lys		
	820	825
		830
Asn Val Val Thr His Arg Leu His Leu Asn Arg Phe Ser Val Ser Gly		
	835	840
		845
Thr Ala Thr Thr Tyr Ser Gln Ser Ser Ala Ser Thr Tyr Val Pro Thr		
	850	855
		860
Val Cys Asn Gly Arg Glu Val Leu Asp Ser Thr Thr Ser Ser Leu		
865	870	875

<210> 54
 <211> 879
 <212> PRT
 <213> Mus musculus

<400> 54

Met Lys Met Leu Thr Arg Leu Gln Val Leu Met Leu Ala Leu Phe Ser
1 5 10 15
Lys Gly Phe Leu Val Ser Leu Gly Asp His Asn Phe Met Arg Arg Glu
20 25 30
Ile Lys Ile Glu Gly Asp Leu Val Leu Gly Gly Leu Phe Pro Ile Asn
35 40 45
Glu Lys Gly Thr Gly Thr Glu Glu Cys Arg Gly Ile Asn Glu Asp Arg
50 55 60
Gly Ile Gln Arg Leu Glu Ala Met Leu Phe Ala Ile Asp Glu Ile Asn
65 70 75 80
Lys Asp Asn Tyr Leu Leu Pro Gly Val Lys Leu Gly Val His Ile Leu
85 90 95
Asp Thr Cys Ser Arg Asp Thr Tyr Ala Leu Glu Gln Ser Leu Glu Phe
100 105 110
Val Arg Ala Ser Leu Thr Lys Val Asp Glu Ala Glu Tyr Met Cys Pro
115 120 125

Asp Gly Ser Tyr Ala Ile Gln Glu Asn Ile Pro Leu Leu Ile Ala Gly
 130 135 140
 Val Ile Gly Gly Ser Tyr Ser Ser Val Ser Ile Gln Val Ala Asn Leu
 145 150 155 160
 Leu Arg Leu Phe Gln Ile Pro Gln Ile Ser Tyr Ala Ser Thr Ser Ala
 165 170 175
 Lys Leu Ser Asp Lys Ser Arg Tyr Asp Tyr Phe Ala Arg Thr Val Pro
 180 185 190
 Pro Asp Phe Tyr Gln Ala Lys Ala Met Ala Glu Ile Leu Arg Tyr Phe
 195 200 205
 Asn Trp Thr Tyr Val Ser Thr Val Ala Ser Glu Gly Asp Tyr Gly Glu
 210 215 220
 Thr Gly Ile Glu Ala Phe Glu Gln Glu Ala Arg Leu Arg Asn Ile Cys
 225 230 235 240
 Ile Ala Thr Ala Glu Lys Val Gly Arg Ser Asn Ile Arg Lys Ser Tyr
 245 250 255
 Asp Ser Val Ile Arg Glu Leu Leu Gln Lys Pro Asn Ala Arg Val Val
 260 265 270
 Val Leu Phe Met Arg Ser Asp Asp Ser Arg Glu Leu Ile Ala Ala Ala
 275 280 285
 Ser Arg Val Asn Ala Ser Phe Thr Trp Val Ala Ser Asp Gly Trp Gly
 290 295 300
 Ala Gln Glu Ser Ile Val Lys Gly Ser Glu His Val Ala Tyr Gly Ala
 305 310 315 320
 Ile Thr Leu Glu Leu Ala Ser His Pro Val Arg Gln Phe Asp Arg Tyr
 325 330 335
 Phe Gln Ser Leu Asn Pro Tyr Asn Asn His Arg Asn Pro Trp Phe Arg
 340 345 350
 Asp Phe Trp Glu Gln Lys Phe Gln Cys Ser Leu Gln Asn Lys Arg Asn
 355 360 365
 His Arg Gln Ile Cys Asp Lys His Leu Ala Ile Asp Ser Ser Asn Tyr
 370 375 380
 Glu Gln Glu Ser Lys Ile Met Phe Val Val Asn Ala Val Tyr Ala Met
 385 390 395 400
 Ala His Ala Leu His Lys Met Gln Arg Thr Leu Cys Pro Asn Thr Thr
 405 410 415
 Lys Leu Cys Asp Ala Met Lys Ile Leu Asp Gly Lys Lys Leu Tyr Lys
 420 425 430

Asp Tyr Leu Leu Lys Ile Asn Phe Thr Ala Pro Phe Asn Pro Asn Lys
 435 440 445
 Gly Ala Asp Ser Ile Val Lys Phe Asp Thr Tyr Gly Asp Gly Met Gly
 450 455 460
 Arg Tyr Asn Val Phe Asn Phe Gln His Ile Gly Gly Lys Tyr Ser Tyr
 465 470 475 480
 Leu Lys Val Gly His Trp Ala Glu Thr Leu Tyr Leu Asp Val Asp Ser
 485 490 495
 Ile His Trp Ser Arg Asn Ser Val Pro Thr Ser Gln Cys Ser Asp Pro
 500 505 510
 Cys Ala Pro Asn Glu Met Lys Asn Met Gln Pro Gly Asp Val Cys Cys
 515 520 525
 Trp Ile Cys Ile Pro Cys Glu Pro Tyr Glu Tyr Leu Val Asp Glu Phe
 530 535 540
 Thr Cys Met Asp Cys Gly Pro Gly Gln Trp Pro Thr Ala Asp Leu Ser
 545 550 555 560
 Gly Cys Tyr Asn Leu Pro Glu Asp Tyr Ile Arg Trp Glu Asp Ala Trp
 565 570 575
 Ala Ile Gly Pro Val Thr Ile Ala Cys Leu Gly Phe Met Cys Thr Cys
 580 585 590
 Ile Val Ile Thr Val Phe Ile Lys His Asn Asn Thr Pro Leu Val Lys
 595 600 605
 Ala Ser Gly Arg Glu Leu Cys Tyr Ile Leu Leu Phe Gly Val Ser Leu
 610 615 620
 Ser Tyr Cys Met Thr Phe Phe Phe Ile Ala Lys Pro Ser Pro Val Ile
 625 630 635 640
 Cys Ala Leu Arg Arg Leu Gly Leu Gly Thr Ser Phe Ala Ile Cys Tyr
 645 650 655
 Ser Ala Leu Leu Thr Lys Thr Asn Cys Ile Ala Arg Ile Phe Asp Gly
 660 665 670
 Val Lys Asn Gly Ala Gln Arg Pro Lys Phe Ile Ser Pro Ser Ser Gln
 675 680 685
 Val Phe Ile Cys Leu Gly Leu Ile Leu Val Gln Ile Val Met Val Ser
 690 695 700
 Val Trp Leu Ile Leu Glu Thr Pro Gly Thr Arg Arg Tyr Thr Leu Pro
 705 710 715 720
 Glu Lys Arg Glu Thr Val Ile Leu Lys Cys Asn Val Lys Asp Ser Ser
 725 730 735

Met Leu Ile Ser Leu Thr Tyr Asp Val Val Leu Val Ile Leu Cys Thr
740 745 750

Val Tyr Ala Phe Lys Thr Arg Lys Cys Pro Glu Asn Phe Asn Glu Ala
755 760 765

Lys Phe Ile Gly Phe Thr Met Tyr Thr Thr Cys Ile Ile Trp Leu Ala
770 775 780

Phe Leu Pro Ile Phe Tyr Val Thr Ser Ser Asp Tyr Arg Val Gln Thr
785 790 795 800

Thr Thr Met Cys Ile Ser Val Ser Leu Ser Gly Phe Val Val Leu Gly
805 810 815

Cys Leu Phe Ala Pro Lys Val His Ile Val Leu Phe Gln Pro Gln Lys
820 825 830

Asn Val Val Thr His Arg Leu His Leu Asn Arg Phe Ser Val Ser Gly
835 840 845

Thr Ala Thr Thr Tyr Ser Gln Ser Ser Ala Ser Thr Tyr Val Pro Thr
850 855 860

Val Cys Asn Gly Arg Glu Val Leu Asp Ser Thr Thr Ser Ser Leu
865 870 875

<210> 55

<211> 442

<212> PRT

<213> Homo sapiens

<400> 55

Met Gly Leu Ala Met Glu His Gly Gly Ser Tyr Ala Arg Ala Gly Gly
1 5 10 15

Ser Ser Arg Gly Cys Trp Tyr Tyr Leu Arg Tyr Phe Phe Leu Phe Val
20 25 30

Ser Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Val
35 40 45

Tyr Gly Asn Val His Val Ser Thr Glu Ser Asn Leu Gln Ala Thr Glu
50 55 60

Arg Arg Ala Glu Gly Leu Tyr Ser Gln Leu Leu Gly Leu Thr Ala Ser
65 70 75 80

Gln Ser Asn Leu Thr Lys Glu Leu Asn Phe Thr Thr Arg Ala Lys Asp
85 90 95

Ala Ile Met Gln Met Trp Leu Asn Ala Arg Arg Asp Leu Asp Arg Ile
100 105 110

Asn Ala Ser Phe Arg Gln Cys Gln Gly Asp Arg Val Ile Tyr Thr Asn
115 120 125

Asn His Arg Tyr Met Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Arg
 130 135 140

Asp Gln Phe Lys Asp Met Asn Lys Ser Cys Asp Ala Leu Leu Phe Met
 145 150 155 160

Leu Asn Gln Lys Val Lys Thr Leu Glu Val Glu Ile Ala Lys Glu Lys
 165 170 175

Thr Ile Cys Thr Lys Asp Lys Glu Ser Val Leu Leu Asn Lys Arg Val
 180 185 190

Ala Glu Glu Gln Leu Val Glu Cys Val Lys Thr Arg Glu Leu Gln His
 195 200 205

Gln Glu Arg Gln Leu Ala Lys Glu Gln Leu Gln Lys Val Gln Ala Leu
 210 215 220

Cys Leu Pro Leu Asp Lys Asp Lys Phe Glu Met Asp Leu Arg Asn Leu
 225 230 235 240

Trp Arg Asp Ser Ile Ile Pro Arg Ser Leu Asp Asn Leu Gly Tyr Asn
 245 250 255

Leu Tyr His Pro Leu Gly Ser Glu Leu Ala Ser Ile Arg Arg Ala Cys
 260 265 270

Asp His Met Pro Ser Leu Met Ser Ser Lys Val Glu Glu Leu Ala Arg
 275 280 285

Ser Leu Arg Ala Asp Ile Glu Arg Val Ala Arg Glu Asn Ser Asp Leu
 290 295 300

Gln Arg Gln Lys Leu Glu Ala Gln Gln Gly Leu Arg Ala Ser Gln Glu
 305 310 315 320

Ala Lys Gln Lys Val Glu Lys Glu Ala Gln Ala Arg Glu Ala Lys Leu
 325 330 335

Gln Ala Glu Cys Ser Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala
 340 345 350

Val Leu Arg Lys Glu Arg Asp Asn Leu Ala Lys Glu Leu Glu Glu Lys
 355 360 365

Lys Arg Glu Ala Glu Gln Leu Arg Met Glu Leu Ala Ile Arg Asn Ser
 370 375 380

Ala Leu Asp Thr Cys Ile Lys Thr Lys Ser Gln Pro Met Met Pro Val
 385 390 395 400

Ser Arg Pro Met Gly Pro Val Pro Asn Pro Gln Pro Ile Asp Pro Ala
 405 410 415

Ser Leu Glu Glu Phe Lys Arg Lys Ile Leu Glu Ser Gln Arg Pro Pro
 420 425 430

Ala Gly Ile Pro Val Ala Pro Ser Ser Gly
 435 440

<210> 56
 <211> 442
 <212> PRT
 <213> Homo sapiens

<400> 56
 Met Gly Leu Ala Met Glu His Gly Gly Ser Tyr Ala Arg Ala Gly Gly
 1 5 10 15
 Ser Ser Arg Gly Cys Trp Tyr Tyr Leu Arg Tyr Phe Phe Leu Phe Val
 20 25 30
 Ser Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Val
 35 40 45
 Tyr Gly Asn Val His Val Ser Thr Glu Ser Asn Leu Gln Ala Thr Glu
 50 55 60
 Arg Arg Ala Glu Gly Leu Tyr Ser Gln Leu Leu Gly Leu Thr Ala Ser
 65 70 75 80
 Gln Ser Asn Leu Thr Lys Glu Leu Asn Phe Thr Thr Arg Ala Lys Asp
 85 90 95
 Ala Ile Met Gln Met Trp Leu Asn Ala Arg Arg Asp Leu Asp Arg Ile
 100 105 110
 Asn Ala Ser Phe Arg Gln Cys Gln Gly Asp Arg Val Ile Tyr Thr Asn
 115 120 125
 Asn Gln Arg Tyr Met Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Arg
 130 135 140
 Asp Gln Phe Lys Asp Met Asn Lys Ser Cys Asp Ala Leu Leu Phe Met
 145 150 155 160
 Leu Asn Gln Lys Val Lys Thr Leu Glu Val Glu Ile Ala Lys Glu Lys
 165 170 175
 Thr Ile Cys Thr Lys Asp Lys Glu Ser Val Leu Leu Asn Lys Arg Val
 180 185 190
 Ala Glu Glu Gln Leu Val Glu Cys Val Lys Thr Arg Glu Leu Gln His
 195 200 205
 Gln Glu Arg Gln Leu Ala Lys Glu Gln Leu Gln Lys Val Gln Ala Leu
 210 215 220
 Cys Leu Pro Leu Asp Lys Asp Lys Phe Glu Met Asp Leu Arg Asn Leu
 225 230 235 240
 Trp Arg Asp Ser Ile Ile Pro Arg Ser Leu Asp Asn Leu Gly Tyr Asn

	245		250		255
Leu Tyr His	Pro Leu Gly Ser Glu	Leu Ala Ser Ile Arg	Arg Ala Cys		
	260		265		270
Asp His Met	Pro Ser Leu Met Ser	Ser Lys Val Glu Glu	Leu Ala Arg		
	275		280		285
Ser Leu Arg	Ala Asp Ile Glu Arg	Val Ala Arg Glu	Asn Ser Asp Leu		
	290		295		300
Gln Arg Gln	Lys Leu Glu Ala Gln	Gln Gly Leu Arg	Ala Ser Gln Glu		
305		310		315	320
Ala Lys Gln	Lys Val Glu Lys Glu	Ala Gln Ala Arg	Glu Ala Lys Leu		
	325		330		335
Gln Ala Glu	Cys Ser Arg Gln Thr	Gln Leu Ala Leu	Glu Glu Lys Ala		
	340		345		350
Val Leu Arg	Lys Glu Arg Asp Asn	Leu Ala Lys Glu	Leu Glu Glu Lys		
	355		360		365
Lys Arg Glu	Ala Glu Gln Leu Arg	Met Glu Leu Ala	Ile Arg Asn Ser		
	370		375		380
Ala Leu Asp	Thr Cys Ile Lys Thr	Lys Ser Gln Pro	Met Met Pro Val		
385		390		395	400
Ser Arg Pro	Met Gly Pro Val Pro	Asn Pro Gln Pro	Ile Asp Pro Ala		
	405		410		415
Ser Leu Glu	Glu Phe Lys Arg Lys	Ile Leu Glu Ser	Gln Arg Pro Pro		
	420		425		430
Ala Gly Ile	Pro Val Ala Pro Ser	Ser Gly			
	435		440		

<210> 57

<211> 438

<212> PRT

<213> Rattus norvegicus

<400> 57

Met Gly Leu	Ser Met Asp Arg	Ser Pro Tyr Ser	Arg Thr Gly Asp	Arg
1		5		10
Asp Arg Gly	Cys Trp Tyr Tyr	Leu Arg Tyr Phe	Phe Leu Phe Val	Ser
	20		25	30
Leu Ile Gln	Phe Leu Ile Ile	Leu Gly Leu Val	Leu Phe Met Ile	Tyr
	35		40	45
Gly Asn Val	His Ala Thr Thr	Glu Ser Ser Leu	Arg Ala Thr Glu	Ile
	50		55	60

Arg Ala Asp Asn Leu Tyr Ser Gln Val Val Gly Leu Ser Ala Ala Gln
 65 70 75 80
 Ala Asn Leu Ser Lys Gln Leu Asn Ile Ser Thr Leu Val Lys Asp Thr
 85 90 95
 Val Met Gln Gln Leu Leu Thr Thr Arg Arg Glu Val Glu Arg Ile Asn
 100 105 110
 Ala Ser Phe Arg Gln Cys Gln Gly Asp Leu Ile Thr Tyr Ile Asn Tyr
 115 120 125
 Asn Arg Phe Ile Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Gln Glu
 130 135 140
 Gln Leu Lys Glu Gly Asn Lys Thr Cys Glu Ala Leu Leu Phe Lys Leu
 145 150 155 160
 Gly Glu Lys Val Lys Thr Leu Glu Met Glu Val Val Lys Glu Lys Ala
 165 170 175
 Val Cys Ser Lys Asp Lys Asp Ser Leu Leu Ala Gly Lys Arg Gln Ala
 180 185 190
 Glu Met Gln Gln Glu Ala Cys Gly Lys Ala Arg Glu Gln Gln Lys Gln
 195 200 205
 Asp Gln Gln Val Thr Glu Glu Gln Leu Arg Lys Val Gln Ser Leu Cys
 210 215 220
 Leu Pro Leu Asp Gln Glu Lys Phe Gln Ala Asp Val Leu Asn Val Trp
 225 230 235 240
 Arg Asp Ser Leu Val Tyr Arg Ser Leu Asp Asn Ile Gly Tyr His Tyr
 245 250 255
 Ser Leu Met Pro Glu Phe Ser Ser Leu Arg Arg Thr Cys Glu Ser Leu
 260 265 270
 Pro Gly Ile Met Thr Thr Lys Val Glu Glu Leu Ala Arg Gly Leu Arg
 275 280 285
 Ala Gly Ile Glu Arg Val Thr Arg Glu Asn Gly Glu Leu Arg Arg Gln
 290 295 300
 Lys Leu Glu Leu Glu Arg Ala Ile Gln Gly Glu Arg Glu Ala Arg Thr
 305 310 315 320
 Arg Ala Gly Thr Glu Ala Gln Ala Arg Glu Thr Gln Leu Arg Thr Glu
 325 330 335
 Cys Ala Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala Ala Leu Arg
 340 345 350
 Thr Gln Arg Asp Asp Leu Glu Arg Gln Leu Glu Ala Arg Lys Arg Glu
 355 360 365

Leu Glu Gln Leu Arg Thr Glu Val Asp Val Arg Ile Ser Ala Leu Asp
 370 375 380
 Thr Cys Val Lys Ala Lys Ser Leu Pro Ala Ile Gln Pro Arg Leu Pro
 385 390 395 400
 Gly Pro Pro Pro Asn Pro Pro Pro Ile Asp Pro Ala Ser Leu Glu Glu
 405 410 415
 Phe Lys Lys Arg Ile Leu Glu Ser Gln Arg Pro Pro Leu Val Asn Pro
 420 425 430
 Ala Val Pro Pro Ser Gly
 435

<210> 58
 <211> 438
 <212> PRT
 <213> Mus musculus

<400> 58
 Met Gly Leu Ser Met Asp Arg Ser Pro Tyr Ala Arg Thr Gly Asp Gln
 1 5 10 15
 Gln Arg Gly Cys Trp Tyr Tyr Leu Arg Tyr Phe Phe Leu Phe Val Ser
 20 25 30
 Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Ile Tyr
 35 40 45
 Gly Asn Val His Ala Thr Thr Glu Ser Ser Leu Arg Ala Thr Glu Ile
 50 55 60
 Arg Ala Asp Ser Leu Tyr Ser Gln Val Val Gly Leu Ser Ala Ser Gln
 65 70 75 80
 Ala Asn Leu Ser Lys Gln Leu Asn Ile Ser Leu Leu Val Lys Glu Thr
 85 90 95
 Val Met Gln Gln Leu Leu Thr Thr Arg Arg Glu Met Glu Arg Ile Asn
 100 105 110
 Ala Ser Phe Arg Gln Cys Gln Gly Asp Leu Ile Thr Tyr Ile Asn Tyr
 115 120 125
 Asn Arg Phe Ile Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Gln Glu
 130 135 140
 Gln Leu Lys Glu Val Asn Lys Thr Cys Glu Ala Leu Leu Phe Lys Leu
 145 150 155 160
 Gly Glu Lys Val Lys Thr Leu Glu Met Glu Val Ala Lys Glu Lys Ala
 165 170 175
 Val Cys Ser Lys Asp Lys Glu Ser Leu Leu Ala Gly Lys Arg Gln Thr
 180 185 190

Glu Glu Gln Leu Glu Ala Cys Gly Lys Ala Arg Glu Arg Gln Gln Gln
 195 200 205
 Glu Gln Gln Val Thr Glu Glu Asn Leu Arg Lys Val Gln Ser Leu Cys
 210 215 220
 Ile Pro Leu Asp Gln Glu Lys Phe Gln Ala Asp Val Leu Ser Ala Trp
 225 230 235 240
 Arg Asp Ser Leu Ile Tyr Arg Thr Leu Glu Thr Leu Pro Tyr His Tyr
 245 250 255
 Gln Leu Met Pro Glu Tyr Ala Ser Leu Arg Arg Thr Cys Glu Ser Leu
 260 265 270
 Pro Gly Ile Met Thr Thr Lys Ile Glu Glu Leu Ala Arg Gly Leu Arg
 275 280 285
 Ala Gly Ile Glu Arg Val Thr Arg Glu Asn Ala Glu Leu Arg Arg Gln
 290 295 300
 Lys Leu Glu Leu Glu Arg Ala Ala Gln Ala Ala Gln Glu Ala Arg Ala
 305 310 315 320
 Arg Ala Gly Thr Glu Ala Gln Ala Arg Glu Thr Gln Leu Arg Ala Glu
 325 330 335
 Cys Ala Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala Ala Leu Arg
 340 345 350
 Ala Gln Arg Asp Asn Leu Glu Arg Glu Leu Glu Ala Arg Lys Arg Glu
 355 360 365
 Leu Glu Gln Leu Arg Thr Glu Val Asp Val Arg Ile Ser Ala Leu Asp
 370 375 380
 Thr Cys Val Lys Ala Lys Ser Leu Pro Ala Val Pro Pro Arg Val Ser
 385 390 395 400
 Gly Pro Pro Pro Asn Pro Pro Pro Ile Asp Pro Ala Ser Leu Glu Glu
 405 410 415
 Phe Lys Lys Arg Ile Leu Glu Ser Gln Arg Leu Pro Val Val Asn Pro
 420 425 430
 Ala Ala Gln Pro Ser Gly
 435

<210> 59

<211> 438

<212> PRT

<213> Mus musculus

<400> 59

Met Gly Leu Ser Met Asp Arg Ser Pro Tyr Ala Arg Thr Gly Asp Gln

1	5	10	15
Gln Arg Gly Cys Trp Tyr Tyr Leu Arg Tyr Phe Phe Leu Phe Val Ser	20	25	30
Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Ile Tyr	35	40	45
Gly Asn Val His Ala Thr Thr Glu Ser Ser Leu Arg Ala Thr Glu Ile	50	55	60
Arg Ala Asp Ser Leu Tyr Ser Gln Val Val Gly Leu Ser Ala Ser Gln	65	70	75
Ala Asn Leu Ser Lys Gln Leu Asn Ile Ser Leu Leu Val Lys Glu Thr	85	90	95
Val Met Gln Gln Leu Leu Thr Thr Arg Arg Glu Met Glu Arg Ile Asn	100	105	110
Ala Ser Phe Arg Gln Cys Gln Gly Asp Leu Ile Thr Tyr Ile Asn Tyr	115	120	125
Asn Arg Phe Ile Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Gln Glu	130	135	140
Gln Leu Lys Glu Val Asn Lys Thr Cys Glu Ala Leu Leu Phe Lys Leu	145	150	155
Gly Glu Lys Val Lys Thr Leu Glu Met Glu Val Ala Lys Glu Lys Ala	165	170	175
Val Cys Ser Lys Asp Lys Glu Ser Leu Leu Ala Gly Lys Arg Gln Ala	180	185	190
Glu Glu Gln Leu Glu Ala Cys Gly Lys Ala Arg Glu Arg Gln Gln Gln	195	200	205
Glu Gln Gln Val Thr Glu Glu Asn Leu Arg Lys Val Gln Ser Leu Cys	210	215	220
Ile Pro Leu Asp Gln Glu Lys Phe Gln Ala Asp Val Leu Ser Ala Trp	225	230	235
Arg Asp Ser Leu Ile Tyr Arg Thr Leu Glu Thr Leu Pro Tyr His Tyr	245	250	255
Gln Leu Met Pro Glu Tyr Ala Ser Leu Arg Arg Thr Cys Glu Ser Leu	260	265	270
Pro Gly Ile Met Pro Pro Lys Ile Glu Glu Met Ala Arg Gly Val Arg	275	280	285
Ala Gly Ile Glu Arg Val Thr Arg Glu Asn Ala Glu Leu Arg Arg Gln	290	295	300
Lys Leu Glu Leu Glu Arg Ala Ala Gln Arg Ala Gln Glu Ala Arg Ala			

305 310 315 320
 Arg Ala Gly Thr Glu Ala Gln Ala Arg Glu Thr Gln Leu Arg Ala Glu
 325 330 335
 Cys Ala Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala Ala Leu Arg
 340 345 350
 Ala Gln Arg Asp Asn Leu Glu Arg Glu Leu Glu Ala Arg Lys Arg Glu
 355 360 365
 Leu Glu Gln Leu Arg Thr Glu Val Asp Val Arg Ile Ser Ala Leu Asp
 370 375 380
 Thr Cys Val Lys Ala Lys Ser Leu Pro Ala Val Pro Pro Arg Val Ser
 385 390 395 400
 Gly Pro Pro Pro Asn Pro Pro Pro Ile Asp Pro Ala Ser Leu Glu Glu
 405 410 415
 Phe Lys Lys Arg Ile Leu Glu Ser Gln Arg Leu Pro Val Val Asn Pro
 420 425 430
 Ala Ala Gln Pro Ser Gly
 435

<210> 60
 <211> 1788
 <212> PRT
 <213> Rattus norvegicus

<400> 60
 Met Leu Arg Pro Met Gln Pro Thr Ser Arg Thr Leu Gln Arg Pro Pro
 1 5 10 15
 Arg Gly Ala Leu Glu Ala Gly Gly Arg Arg Asn Cys Gln Asp Gln Val
 20 25 30
 Ala His Pro Asn Trp Asn Thr Gln Ser Val Gln Thr Pro Arg Val Arg
 35 40 45
 Arg Thr Leu Gly Ala Pro Val Pro Pro Ser Arg Lys Val Lys Ala Trp
 50 55 60
 Ala Pro Gly Thr Asp Gln Trp Pro Gly Val Ser Pro His Cys Lys Arg
 65 70 75 80
 Ser Glu Ala Glu Ala Lys Pro Ser Gly Ser Gln Thr Val Asn Leu Thr
 85 90 95
 Gly Arg Ala Asn Asp Pro Cys Asp Leu Asp Ser Arg Val Gln Ala Thr
 100 105 110
 Ser Val Lys Val Thr Val Ala Gly Phe Gln Pro Gly Gly Ala Val Glu
 115 120 125

Lys Leu Cys Gln Glu Ser Leu Gly Lys Leu Thr Thr Gly Asp Ala Cys
 130 135 140

Val Ser Thr Ser Cys Glu Leu Ala Ser Ala Leu Ser His Leu Asp Ala
 145 150 155 160

Ser His Leu Thr Glu Asn Leu Pro Lys Ala Ala Ser Glu Leu Gly Gln
 165 170 175

Gln Pro Met Thr Ser Ser Asp Leu Ile Ser Ser Pro Gly Lys Lys Gly
 180 185 190

Ala Ala His Pro Asp Pro Ser Lys Thr Ser Val Asp Thr Gly Gln Val
 195 200 205

Ser Arg Pro Glu Asn Pro Ser Gln Pro Ala Ser Pro Arg Val Thr Lys
 210 215 220

Cys Lys Ala Arg Ser Pro Val Arg Leu Pro His Glu Gly Ser Pro Ser
 225 230 235 240

Pro Gly Glu Lys Ala Ala Ala Pro Pro Asp Tyr Ser Lys Thr Arg Ser
 245 250 255

Ala Ser Glu Thr Ser Thr Pro His Asn Thr Arg Arg Val Ala Ala Leu
 260 265 270

Arg Gly Ala Gly Pro Gly Ala Glu Gly Met Thr Pro Ala Gly Ala Val
 275 280 285

Leu Pro Gly Asp Pro Leu Thr Ser Gln Glu Gln Arg Gln Gly Ala Pro
 290 295 300

Gly Asn His Ser Lys Ala Leu Glu Met Thr Gly Ile His Ala Pro Glu
 305 310 315 320

Ser Ser Gln Glu Pro Ser Leu Leu Glu Gly Ala Asp Ser Val Ser Ser
 325 330 335

Arg Ala Pro Gln Ala Ser Leu Ser Met Leu Pro Ser Thr Asp Asn Thr
 340 345 350

Lys Glu Ala Cys Gly His Val Ser Gly His Cys Cys Pro Gly Gly Ser
 355 360 365

Arg Glu Ser Pro Val Thr Asp Ile Asp Ser Phe Ile Lys Glu Leu Asp
 370 375 380

Ala Ser Ala Ala Arg Ser Pro Ser Ser Gln Thr Gly Asp Ser Gly Ser
 385 390 395 400

Gln Glu Gly Ser Ala Gln Gly His Pro Pro Ala Gly Ala Gly Gly Gly
 405 410 415

Ser Ser Cys Arg Ala Glu Pro Val Pro Gly Gly Gln Thr Ser Ser Pro
 420 425 430

Arg Arg Ala Trp Ala Ala Gly Ala Pro Ala Tyr Pro Gln Trp Ala Ser
 435 440 445
 Gln Pro Ser Val Leu Asp Ser Ile Asn Pro Asp Lys His Phe Thr Val
 450 455 460
 Asn Lys Asn Phe Leu Ser Asn Tyr Ser Arg Asn Phe Ser Ser Phe His
 465 470 475 480
 Glu Asp Ser Thr Ser Leu Ser Gly Leu Gly Asp Ser Thr Glu Pro Ser
 485 490 495
 Leu Ser Ser Met Tyr Gly Asp Ala Glu Asp Ser Ser Ser Asp Pro Glu
 500 505 510
 Ser Leu Thr Glu Ala Pro Arg Ala Ser Ala Arg Asp Gly Trp Ser Pro
 515 520 525
 Pro Arg Ser Arg Val Ser Leu His Lys Glu Asp Pro Ser Glu Ser Glu
 530 535 540
 Glu Glu Gln Ile Glu Ile Cys Ser Thr Arg Gly Cys Pro Asn Pro Pro
 545 550 555 560
 Ser Ser Pro Ala His Leu Pro Thr Gln Ala Ala Ile Cys Pro Ala Ser
 565 570 575
 Ala Lys Val Leu Ser Leu Lys Tyr Ser Thr Pro Arg Glu Ser Val Ala
 580 585 590
 Ser Pro Arg Glu Lys Ala Ala Cys Leu Pro Gly Ser Tyr Thr Ser Gly
 595 600 605
 Pro Asp Ser Ser Gln Pro Ser Ser Leu Leu Glu Met Ser Ser Gln Glu
 610 615 620
 His Glu Thr His Ala Asp Ile Ser Thr Ser Gln Asn His Arg Pro Ser
 625 630 635 640
 Cys Ala Glu Glu Thr Thr Glu Val Thr Ser Ala Ser Ser Ala Met Glu
 645 650 655
 Asn Ser Pro Leu Ser Lys Val Ala Arg His Phe His Ser Pro Pro Ile
 660 665 670
 Ile Leu Ser Ser Pro Asn Met Val Asn Gly Leu Glu His Asp Leu Leu
 675 680 685
 Asp Asp Glu Thr Leu Asn Gln Tyr Glu Thr Ser Ile Asn Ala Ala Ala
 690 695 700
 Ser Leu Ser Ser Phe Ser Val Asp Val Pro Lys Asn Gly Glu Ser Val
 705 710 715 720
 Leu Glu Asn Leu His Ile Ser Glu Ser Gln Asp Leu Asp Asp Leu Leu
 725 730 735

Gln Lys Pro Lys Met Ile Ala Arg Arg Pro Ile Met Ala Trp Phe Lys
 740 745 750
 Glu Ile Asn Lys His Asn Gln Gly Thr His Leu Arg Ser Lys Thr Glu
 755 760 765
 Lys Glu Gln Pro Leu Met Pro Ala Arg Ser Pro Asp Ser Lys Ile Gln
 770 775 780
 Met Val Ser Ser Ser Gln Lys Lys Gly Val Thr Val Pro His Ser Pro
 785 790 795 800
 Pro Gln Pro Lys Thr Asn Leu Glu Asn Lys Asp Leu Ser Lys Lys Ser
 805 810 815
 Pro Ala Glu Met Leu Leu Thr Asn Gly Gln Lys Ala Lys Cys Gly Pro
 820 825 830
 Lys Leu Lys Arg Leu Ser Leu Lys Gly Lys Ala Lys Val Asn Ser Glu
 835 840 845
 Ala Pro Ala Ala Asn Ala Val Lys Ala Gly Gly Thr Asp His Arg Lys
 850 855 860
 Pro Leu Ile Ser Pro Gln Thr Ser His Lys Thr Leu Ser Lys Ala Val
 865 870 875 880
 Ser Gln Arg Leu His Val Ala Asp His Glu Asp Pro Asp Arg Asn Thr
 885 890 895
 Thr Ala Ala Pro Arg Ser Pro Gln Cys Val Leu Glu Ser Lys Pro Pro
 900 905 910
 Leu Ala Thr Ser Gly Pro Leu Lys Pro Ser Val Ser Asp Thr Ser Ile
 915 920 925
 Arg Thr Phe Val Ser Pro Leu Thr Ser Pro Lys Pro Val Pro Glu Gln
 930 935 940
 Gly Met Trp Ser Arg Phe His Met Ala Val Leu Ser Glu Pro Asp Arg
 945 950 955 960
 Gly Cys Pro Thr Thr Pro Lys Ser Pro Lys Cys Arg Ala Glu Gly Arg
 965 970 975
 Ala Pro Arg Ala Asp Ser Gly Pro Val Ser Pro Ala Ala Ser Arg Asn
 980 985 990
 Gly Met Ser Val Ala Gly Asn Arg Gln Ser Glu Pro Arg Leu Ala Ser
 995 1000 1005
 His Val Ala Ala Asp Thr Ala Gln Pro Arg Pro Thr Gly Glu Lys Gly
 1010 1015 1020
 Gly Asn Ile Met Ala Ser Asp Arg Leu Glu Arg Thr Asn Gln Leu Lys
 1025 1030 1035 1040

Ile Val Glu Ile Ser Ala Glu Ala Val Ser Glu Thr Val Cys Gly Asn
 1045 1050 1055
 Lys Pro Ala Glu Ser Asp Arg Arg Gly Gly Cys Leu Ala Gln Gly Asn
 1060 1065 1070
 Cys Gln Glu Lys Ser Glu Ile Arg Leu Tyr Arg Gln Val Ala Glu Ser
 1075 1080 1085
 Ser Thr Ser His Pro Ser Ser Leu Pro Ser His Ala Ser Gln Ala Glu
 1090 1095 1100
 Gln Glu Met Ser Arg Ser Phe Ser Met Ala Lys Leu Ala Ser Ser Ser
 1105 1110 1115 1120
 Ser Ser Leu Gln Thr Ala Ile Arg Lys Ala Glu Tyr Ser Gln Gly Lys
 1125 1130 1135
 Ser Ser Leu Met Ser Asp Ser Arg Gly Val Pro Arg Asn Ser Ile Pro
 1140 1145 1150
 Gly Gly Pro Ser Gly Glu Asp His Leu Tyr Phe Thr Pro Arg Pro Ala
 1155 1160 1165
 Thr Arg Thr Tyr Ser Met Pro Ala Gln Phe Ser Ser His Phe Gly Arg
 1170 1175 1180
 Glu Gly His Pro Pro His Ser Leu Gly Arg Ser Arg Asp Ser Gln Val
 1185 1190 1195 1200
 Pro Val Thr Ser Ser Val Val Pro Glu Ala Lys Ala Ser Arg Gly Gly
 1205 1210 1215
 Leu Pro Ser Leu Ala Asn Gly Gln Gly Ile Tyr Ser Val Lys Pro Leu
 1220 1225 1230
 Leu Asp Thr Ser Arg Asn Leu Pro Ala Thr Asp Glu Gly Asp Ile Ile
 1235 1240 1245
 Ser Val Gln Glu Thr Ser Cys Leu Val Thr Asp Lys Ile Lys Val Thr
 1250 1255 1260
 Arg Arg His Tyr Cys Tyr Glu Gln Asn Trp Pro His Glu Ser Thr Ser
 1265 1270 1275 1280
 Phe Phe Ser Val Lys Gln Arg Ile Lys Ser Phe Glu Asn Leu Ala Asn
 1285 1290 1295
 Ala Asp Arg Pro Val Ala Lys Ser Gly Ala Ser Pro Phe Leu Ser Val
 1300 1305 1310
 Ser Ser Lys Pro Pro Ile Gly Arg Arg Ser Ser Gly Ser Ile Val Ser
 1315 1320 1325
 Gly Ser Leu Gly His Pro Gly Asp Ala Ala Ala Arg Leu Leu Arg Arg
 1330 1335 1340

Ser Leu Ser Ser Cys Ser Glu Asn Gln Ser Glu Ala Gly Thr Leu Leu
 1345 1350 1355 1360
 Pro Gln Met Ala Lys Ser Pro Ser Ile Met Thr Leu Thr Ile Ser Arg
 1365 1370 1375
 Gln Asn Pro Pro Glu Thr Ser Ser Lys Gly Ser Asp Ser Glu Leu Lys
 1380 1385 1390
 Lys Ser Leu Gly Pro Leu Gly Ile Pro Thr Pro Thr Met Thr Leu Ala
 1395 1400 1405
 Ser Pro Val Lys Arg Asn Lys Ser Ser Val Arg His Thr Gln Pro Ser
 1410 1415 1420
 Pro Val Ser Arg Ser Lys Leu Gln Glu Leu Arg Ala Leu Ser Met Pro
 1425 1430 1435 1440
 Asp Leu Asp Lys Leu Cys Ser Glu Asp Tyr Ser Ala Gly Pro Ser Ala
 1445 1450 1455
 Val Leu Phe Lys Thr Glu Leu Glu Ile Thr Pro Arg Arg Ser Pro Gly
 1460 1465 1470
 Pro Pro Ala Gly Gly Val Ser Cys Pro Glu Lys Gly Gly Asn Arg Ala
 1475 1480 1485
 Cys Pro Gly Gly Ser Gly Pro Lys Thr Ser Ala Ala Glu Thr Pro Ser
 1490 1495 1500
 Ser Ala Ser Asp Thr Gly Glu Ala Ala Gln Asp Leu Pro Phe Arg Arg
 1505 1510 1515 1520
 Ser Trp Ser Val Lys Leu Asp Gln Leu Leu Val Ser Ala Gly Asp Gln
 1525 1530 1535
 Gln Arg Leu Gln Ser Val Leu Ser Ser Val Gly Ser Lys Ser Thr Ile
 1540 1545 1550
 Leu Thr Leu Ile Gln Glu Ala Lys Ala Gln Ser Glu Asn Glu Glu Asp
 1555 1560 1565
 Val Cys Phe Ile Val Leu Asn Arg Lys Glu Gly Ser Gly Leu Gly Phe
 1570 1575 1580
 Ser Val Ala Gly Gly Thr Asp Val Glu Pro Lys Ser Ile Thr Val His
 1585 1590 1595 1600
 Arg Val Phe Ser Gln Gly Ala Ala Ser Gln Glu Gly Thr Met Asn Arg
 1605 1610 1615
 Gly Asp Phe Leu Leu Ser Val Asn Gly Ala Ser Leu Ala Gly Leu Ala
 1620 1625 1630
 His Gly Asn Val Leu Lys Val Leu His Gln Ala Gln Leu His Lys Asp
 1635 1640 1645

Ala Leu Val Val Ile Lys Lys Gly Met Asp Gln Pro Arg Pro Ser Ala
 1650 1655 1660

Arg Gln Glu Pro Pro Thr Ala Asn Gly Lys Gly Leu Leu Ser Arg Lys
 1665 1670 1675 1680

Thr Ile Pro Leu Glu Pro Gly Ile Gly Arg Ser Val Ala Val His Asp
 1685 1690 1695

Ala Leu Cys Val Glu Val Leu Lys Thr Ser Ala Gly Leu Gly Leu Ser
 1700 1705 1710

Leu Asp Gly Gly Lys Ser Ser Val Thr Gly Asp Gly Pro Leu Val Ile
 1715 1720 1725

Lys Arg Val Tyr Lys Gly Gly Ala Ala Glu Gln Ala Gly Ile Ile Glu
 1730 1735 1740

Ala Gly Asp Glu Ile Leu Ala Ile Asn Gly Lys Pro Leu Val Gly Leu
 1745 1750 1755 1760

Met His Phe Asp Ala Trp Asn Ile Met Lys Ser Val Pro Glu Gly Pro
 1765 1770 1775

Val Gln Leu Leu Ile Arg Lys His Arg Asn Ser Ser
 1780 1785

<210> 61
 <211> 1608
 <212> PRT
 <213> Homo sapiens

<400> 61
 Ser Ser Asp Leu Ile Ser Ser Pro Gly Lys Lys Gly Ala Ala His Pro
 1 5 10 15

Asp Pro Ser Lys Thr Ser Val Asp Thr Gly Lys Val Ser Arg Pro Glu
 20 25 30

Asn Pro Ser Gln Pro Ala Ser Pro Arg Val Ala Lys Cys Lys Ala Arg
 35 40 45

Ser Pro Val Arg Leu Pro His Glu Gly Ser Pro Ser Pro Gly Glu Lys
 50 55 60

Ala Ala Ala Pro Pro Asp Tyr Ser Lys Thr Arg Ser Ala Ser Glu Thr
 65 70 75 80

Ser Thr Pro His Asn Thr Arg Arg Val Ala Ala Leu Arg Gly Ala Gly
 85 90 95

Pro Gly Ala Glu Gly Met Thr Pro Ala Gly Ala Val Leu Pro Gly Asp
 100 105 110

Pro Leu Thr Ser Gln Glu Gln Arg Gln Gly Ala Pro Gly Asn His Ser
 115 120 125

Lys Ala Leu Glu Met Thr Gly Ile His Ala Pro Glu Ser Ser Gln Glu
 130 135 140
 Pro Ser Leu Leu Glu Gly Ala Asp Ser Val Ser Ser Arg Ala Pro Gln
 145 150 155 160
 Ala Ser Leu Ser Met Leu Pro Ser Thr Asp Asn Thr Lys Glu Ala Cys
 165 170 175
 Gly His Val Ser Gly His Cys Cys Pro Gly Gly Ser Arg Glu Ser Pro
 180 185 190
 Val Thr Asp Ile Asp Ser Phe Ile Lys Glu Leu Asp Ala Ser Ala Ala
 195 200 205
 Arg Ser Pro Ser Ser Gln Thr Gly Asp Ser Gly Ser Gln Glu Gly Ser
 210 215 220
 Ala Gln Gly His Pro Pro Ala Gly Ala Gly Gly Gly Ser Ser Cys Arg
 225 230 235 240
 Ala Glu Pro Val Pro Gly Gly Gln Thr Ser Ser Pro Arg Arg Ala Trp
 245 250 255
 Ala Ala Gly Ala Pro Ala Tyr Pro Gln Trp Ala Ser Gln Pro Ser Val
 260 265 270
 Leu Asp Ser Ile Asn Pro Asp Lys His Phe Thr Val Asn Lys Asn Phe
 275 280 285
 Leu Ser Asn Tyr Ser Arg Asn Phe Ser Ser Phe His Glu Asp Ser Thr
 290 295 300
 Ser Leu Ser Gly Leu Gly Asp Ser Thr Glu Pro Ser Leu Ser Ser Met
 305 310 315 320
 Tyr Gly Asp Ala Glu Asp Ser Ser Ser Asp Pro Glu Ser Leu Thr Glu
 325 330 335
 Ala Pro Arg Ala Ser Ala Arg Asp Gly Trp Ser Pro Pro Arg Ser Arg
 340 345 350
 Val Ser Leu His Lys Glu Asp Pro Ser Glu Ser Glu Glu Glu Gln Ile
 355 360 365
 Glu Ile Cys Ser Thr Arg Gly Cys Pro Asn Pro Pro Ser Ser Pro Ala
 370 375 380
 His Leu Pro Thr Gln Ala Ala Ile Cys Pro Ala Ser Ala Lys Val Leu
 385 390 395 400
 Ser Leu Lys Tyr Ser Thr Pro Arg Glu Ser Val Ala Ser Pro Arg Glu
 405 410 415
 Lys Val Ala Cys Leu Pro Gly Ser Tyr Thr Ser Gly Pro Asp Ser Ser
 420 425 430

Gln	Pro	Ser	Ser	Leu	Leu	Glu	Met	Ser	Ser	Gln	Glu	His	Glu	Thr	His	
		435					440					445				
Ala	Asp	Ile	Ser	Thr	Ser	Gln	Asn	His	Arg	Pro	Ser	Cys	Ala	Glu	Glu	
	450					455					460					
Thr	Thr	Glu	Val	Thr	Ser	Ala	Ser	Ser	Ala	Met	Glu	Asn	Ser	Pro	Leu	
465					470					475					480	
Ser	Lys	Val	Ala	Arg	His	Phe	His	Ser	Pro	Pro	Ile	Ile	Leu	Ser	Ser	
				485					490					495		
Pro	Asn	Met	Val	Asn	Gly	Leu	Glu	His	Asp	Leu	Leu	Asp	Asp	Glu	Thr	
			500					505					510			
Leu	Asn	Gln	Tyr	Glu	Thr	Ser	Ile	Asn	Ala	Ala	Ala	Ser	Leu	Ser	Ser	
		515					520					525				
Phe	Ser	Val	Asp	Val	Pro	Lys	Asn	Gly	Glu	Ser	Val	Leu	Glu	Asn	Leu	
	530					535					540					
His	Ile	Ser	Glu	Ser	Gln	Asp	Leu	Asp	Asp	Leu	Leu	Gln	Lys	Pro	Lys	
545					550					555					560	
Met	Ile	Ala	Arg	Arg	Pro	Ile	Met	Ala	Trp	Phe	Lys	Glu	Ile	Asn	Lys	
				565					570					575		
His	Asn	Gln	Gly	Thr	His	Leu	Arg	Ser	Lys	Thr	Glu	Lys	Glu	Gln	Pro	
			580					585					590			
Leu	Met	Pro	Ala	Arg	Ser	Pro	Asp	Ser	Lys	Ile	Gln	Met	Val	Ser	Ser	
		595					600					605				
Ser	Gln	Lys	Lys	Gly	Val	Thr	Val	Pro	His	Ser	Pro	Pro	Gln	Pro	Lys	
	610					615					620					
Thr	Asn	Leu	Glu	Asn	Lys	Asp	Leu	Ser	Lys	Lys	Ser	Pro	Ala	Glu	Met	
625					630					635					640	
Leu	Leu	Thr	Asn	Gly	Gln	Lys	Ala	Lys	Cys	Gly	Pro	Lys	Leu	Lys	Arg	
				645					650					655		
Leu	Ser	Leu	Lys	Gly	Lys	Ala	Lys	Val	Asn	Ser	Glu	Ala	Pro	Ala	Ala	
			660					665					670			
Asn	Ala	Val	Lys	Ala	Gly	Gly	Thr	Asp	His	Arg	Lys	Pro	Leu	Ile	Ser	
		675					680					685				
Pro	Gln	Thr	Ser	His	Lys	Thr	Leu	Ser	Lys	Ala	Val	Ser	Gln	Arg	Leu	
		690				695					700					
His	Val	Ala	Asp	His	Glu	Asp	Pro	Asp	Arg	Asn	Thr	Thr	Ala	Ala	Pro	
705					710					715					720	
Arg	Ser	Pro	Gln	Cys	Val	Leu	Glu	Ser	Lys	Pro	Pro	Leu	Ala	Thr	Ser	
				725					730					735		

Gly Pro Leu Lys Pro Ser Val Ser Asp Thr Ser Ile Arg Thr Phe Val
 740 745 750
 Ser Pro Leu Thr Ser Pro Lys Pro Val Pro Glu Gln Gly Met Trp Ser
 755 760 765
 Arg Phe His Met Ala Val Leu Ser Glu Pro Asp Arg Gly Cys Pro Thr
 770 775 780
 Thr Pro Lys Ser Pro Lys Cys Arg Ala Glu Gly Arg Ala Pro Arg Ala
 785 790 795 800
 Asp Ser Gly Pro Val Ser Pro Ala Ala Ser Arg Asn Gly Met Ser Val
 805 810 815
 Ala Gly Asn Arg Gln Ser Glu Pro Arg Leu Ala Ser His Val Ala Ala
 820 825 830
 Asp Thr Ala Gln Pro Arg Pro Thr Gly Glu Lys Gly Gly Asn Ile Met
 835 840 845
 Ala Ser Asp Arg Leu Glu Arg Thr Asn Gln Leu Lys Ile Val Glu Ile
 850 855 860
 Ser Ala Glu Ala Val Ser Glu Thr Val Cys Gly Asn Lys Pro Ala Glu
 865 870 875 880
 Ser Asp Arg Arg Gly Gly Cys Leu Ala Gln Gly Asn Cys Gln Glu Lys
 885 890 895
 Ser Glu Ile Arg Leu Tyr Arg Gln Val Ala Glu Ser Ser Thr Ser His
 900 905 910
 Pro Ser Ser Leu Pro Ser His Ala Ser Gln Ala Glu Gln Glu Met Ser
 915 920 925
 Arg Ser Phe Ser Met Ala Lys Leu Ala Ser Ser Ser Ser Ser Leu Gln
 930 935 940
 Thr Ala Ile Arg Lys Ala Glu Tyr Ser Gln Gly Lys Ser Ser Leu Met
 945 950 955 960
 Ser Asp Ser Arg Gly Val Pro Arg Asn Ser Ile Pro Gly Gly Pro Ser
 965 970 975
 Gly Glu Asp His Leu Tyr Phe Thr Pro Arg Pro Ala Thr Arg Thr Tyr
 980 985 990
 Ser Met Pro Ala Gln Phe Ser Ser His Phe Gly Arg Glu Gly His Pro
 995 1000 1005
 Pro His Ser Leu Gly Arg Ser Arg Asp Ser Gln Val Pro Val Thr Ser
 1010 1015 1020
 Ser Val Val Pro Glu Ala Lys Ala Ser Arg Gly Gly Leu Pro Ser Leu
 1025 1030 1035 1040

Ala Asn Gly Gln Gly Ile Tyr Ser Val Lys Pro Leu Leu Asp Thr Ser
 1045 1050 1055
 Arg Asn Leu Pro Ala Thr Asp Glu Gly Asp Ile Ile Ser Val Gln Glu
 1060 1065 1070
 Thr Ser Cys Leu Val Thr Asp Lys Ile Lys Val Thr Arg Arg His Tyr
 1075 1080 1085
 Cys Tyr Glu Gln Asn Trp Pro His Glu Ser Thr Ser Phe Phe Ser Val
 1090 1095 1100
 Lys Gln Arg Ile Lys Ser Phe Glu Asn Leu Ala Asn Ala Asp Arg Pro
 1105 1110 1115 1120
 Val Ala Lys Ser Gly Ala Ser Pro Phe Leu Ser Val Ser Ser Lys Pro
 1125 1130 1135
 Pro Ile Gly Arg Arg Ser Ser Gly Ser Ile Val Ser Gly Ser Leu Gly
 1140 1145 1150
 His Pro Gly Asp Ala Ala Ala Arg Leu Leu Arg Arg Ser Leu Ser Ser
 1155 1160 1165
 Cys Ser Glu Asn Gln Ser Glu Ala Gly Thr Leu Leu Pro Gln Met Ala
 1170 1175 1180
 Lys Ser Pro Ser Ile Met Thr Leu Thr Ile Ser Arg Gln Asn Pro Pro
 1185 1190 1195 1200
 Glu Thr Ser Ser Lys Gly Ser Asp Ser Glu Leu Lys Lys Ser Leu Gly
 1205 1210 1215
 Pro Leu Gly Ile Pro Thr Pro Thr Met Thr Leu Ala Ser Pro Val Lys
 1220 1225 1230
 Arg Asn Lys Ser Ser Val Arg His Thr Gln Pro Ser Pro Val Ser Arg
 1235 1240 1245
 Ser Lys Leu Gln Glu Leu Arg Ala Leu Ser Met Pro Asp Leu Asp Lys
 1250 1255 1260
 Leu Cys Ser Glu Asp Tyr Ser Ala Gly Pro Ser Ala Val Leu Phe Lys
 1265 1270 1275 1280
 Thr Glu Leu Glu Ile Thr Pro Arg Arg Ser Pro Gly Pro Pro Ala Gly
 1285 1290 1295
 Gly Val Ser Cys Pro Glu Lys Gly Gly Asn Arg Ala Cys Pro Gly Gly
 1300 1305 1310
 Ser Gly Pro Lys Thr Ser Ala Ala Glu Thr Pro Ser Ser Ala Ser Asp
 1315 1320 1325
 Thr Gly Glu Ala Ala Gln Asp Leu Pro Phe Arg Arg Ser Trp Ser Val
 1330 1335 1340

Asn Leu Asp Gln Leu Leu Val Ser Ala Gly Asp Gln Gln Arg Leu Gln
 1345 1350 1355 1360
 Ser Val Leu Ser Ser Val Gly Ser Lys Ser Thr Ile Leu Thr Leu Ile
 1365 1370 1375
 Gln Glu Ala Lys Ala Gln Ser Glu Asn Glu Glu Asp Val Cys Phe Ile
 1380 1385 1390
 Val Leu Asn Arg Lys Glu Gly Ser Gly Leu Gly Phe Ser Val Ala Gly
 1395 1400 1405
 Gly Thr Asp Val Glu Pro Lys Ser Ile Thr Val His Arg Val Phe Ser
 1410 1415 1420
 Gln Gly Ala Ala Ser Gln Glu Gly Thr Met Asn Arg Gly Asp Phe Leu
 1425 1430 1435 1440
 Leu Ser Val Asn Gly Ala Ser Leu Ala Gly Leu Ala His Gly Asn Val
 1445 1450 1455
 Leu Lys Val Leu His Gln Ala Gln Leu His Lys Asp Ala Leu Val Val
 1460 1465 1470
 Ile Lys Lys Gly Met Asp Gln Pro Arg Pro Ser Ala Arg Gln Glu Pro
 1475 1480 1485
 Pro Thr Ala Asn Gly Lys Gly Leu Leu Ser Arg Lys Thr Ile Pro Leu
 1490 1495 1500
 Glu Pro Gly Ile Gly Arg Ser Val Ala Val His Asp Ala Leu Cys Val
 1505 1510 1515 1520
 Glu Val Leu Lys Thr Ser Ala Gly Leu Gly Leu Ser Leu Asp Gly Gly
 1525 1530 1535
 Lys Ser Ser Val Thr Gly Asp Gly Pro Leu Val Ile Lys Arg Val Tyr
 1540 1545 1550
 Lys Gly Gly Ala Ala Glu Gln Ala Gly Ile Ile Glu Ala Gly Asp Glu
 1555 1560 1565
 Ile Leu Ala Ile Asn Gly Lys Pro Leu Val Gly Leu Met His Phe Asp
 1570 1575 1580
 Ala Trp Asn Ile Met Lys Ser Val Pro Glu Gly Pro Val Gln Leu Leu
 1585 1590 1595 1600
 Ile Arg Lys His Arg Asn Ser Ser
 1605

<210> 62

<211> 2766

<212> PRT

<213> Rattus norvegicus

<400> 62

Met	Pro	Ile	Thr	Gln	Asp	Asn	Ala	Leu	Leu	His	Leu	Pro	Leu	Leu	Tyr	1	5	10	15
Glu	Trp	Leu	Gln	Asn	Ser	Leu	Arg	Glu	Gly	Gly	Asp	Ser	Pro	Glu	Gln	20	25	30	
Arg	Leu	Cys	Gln	Ala	Ala	Ile	Gln	Lys	Leu	Gln	Glu	Tyr	Ile	Gln	Leu	35	40	45	
Asn	Leu	Ala	Val	Asp	Glu	Ser	Thr	Val	Pro	Pro	Asp	His	Ser	Pro	Pro	50	55	60	
Glu	Met	Glu	Ile	Cys	Thr	Val	Tyr	Leu	Thr	Lys	Gln	Leu	Gly	Asp	Thr	65	70	75	80
Glu	Thr	Val	Gly	Leu	Ser	Phe	Gly	Asn	Ile	Pro	Val	Phe	Gly	Asp	Tyr	85	90	95	
Gly	Glu	Lys	Arg	Arg	Gly	Gly	Lys	Lys	Arg	Lys	Thr	His	Gln	Gly	Pro	100	105	110	
Val	Leu	Asp	Val	Gly	Cys	Ile	Trp	Val	Thr	Glu	Leu	Arg	Lys	Asn	Ser	115	120	125	
Pro	Ala	Gly	Lys	Ser	Gly	Lys	Val	Arg	Leu	Arg	Asp	Glu	Ile	Leu	Ser	130	135	140	
Leu	Asn	Gly	Gln	Leu	Met	Val	Gly	Val	Asp	Val	Thr	Gly	Ala	Ser	Tyr	145	150	155	160
Leu	Ala	Glu	Gln	Cys	Trp	Asn	Gly	Gly	Phe	Ile	Tyr	Leu	Ile	Met	Leu	165	170	175	
Arg	Arg	Phe	Lys	Gln	Lys	Ala	His	Val	Thr	Tyr	Asn	Gly	Asn	Ser	Gly	180	185	190	
Asn	Ser	Ser	Glu	Pro	Gly	Glu	Thr	Pro	Thr	Leu	Glu	Leu	Gly	Asp	Gln	195	200	205	
Thr	Ser	Lys	Lys	Gly	Lys	Arg	Thr	Arg	Lys	Phe	Gly	Val	Ile	Ser	Arg	210	215	220	
Pro	Ser	Ile	Ser	Lys	Thr	Pro	Glu	Asp	Ser	Lys	Ser	Ser	Ser	Gly	Cys	225	230	235	240
Asp	Thr	Ala	Asp	Asp	Pro	Asn	Ser	Glu	Leu	Glu	Asn	Gly	Ala	Asp	Pro	245	250	255	
Glu	Leu	Gly	Asn	Gly	His	Ala	Phe	Glu	Leu	Glu	Asn	Gly	Pro	His	Ser	260	265	270	
Leu	Lys	Asp	Val	Ala	Gly	Pro	His	Leu	Glu	Arg	Ser	Glu	Ala	Asp	Ser	275	280	285	
Glu	Val	Glu	Leu	Arg	Val	Pro	Lys	Thr	Glu	Ala	Pro	Leu	Ser	Asp	Ser				

290					295					300					
Asn	Asp	Lys	Arg	Arg	Phe	Ser	Lys	Thr	Gly	Lys	Thr	Asp	Phe	Gln	Ser
305					310					315					320
Ser	Asp	Cys	Leu	Ala	Arg	Glu	Glu	Val	Gly	Arg	Ile	Trp	Lys	Met	Glu
			325						330					335	
Leu	Leu	Lys	Glu	Ser	Asp	Gly	Leu	Gly	Ile	Gln	Val	Ser	Gly	Gly	Arg
			340					345					350		
Gly	Ser	Lys	Arg	Ser	Pro	His	Ala	Ile	Val	Val	Thr	Gln	Val	Lys	Glu
		355					360					365			
Gly	Gly	Ala	Ala	His	Arg	Asp	Gly	Arg	Leu	Ser	Leu	Gly	Asp	Glu	Leu
	370					375					380				
Leu	Val	Ile	Asn	Gly	His	Leu	Leu	Val	Gly	Leu	Ser	His	Glu	Glu	Ala
385					390					395					400
Val	Ala	Ile	Leu	Arg	Ser	Ala	Thr	Gly	Met	Val	Gln	Leu	Val	Val	Ala
				405					410					415	
Ser	Lys	Met	Pro	Gly	Ser	Glu	Glu	Ser	Gln	Asp	Val	Gly	Ser	Ser	Glu
			420					425					430		
Glu	Ser	Lys	Gly	Asn	Leu	Glu	Ser	Pro	Lys	Gln	Gly	Asn	Cys	Lys	Thr
		435					440					445			
Lys	Leu	Lys	Ser	Arg	Leu	Ser	Gly	Gly	Val	His	Arg	Leu	Glu	Ser	Val
	450					455					460				
Glu	Glu	Tyr	Asn	Glu	Leu	Met	Val	Arg	Asn	Gly	Asp	Pro	Arg	Ile	Arg
465					470					475					480
Met	Leu	Glu	Val	Ser	Arg	Asp	Gly	Arg	Lys	His	Ser	Leu	Pro	Gln	Leu
				485					490					495	
Leu	Asp	Ser	Thr	Gly	Thr	Ser	Gln	Glu	Tyr	His	Ile	Val	Lys	Lys	Ser
			500					505					510		
Thr	Arg	Ser	Leu	Ser	Thr	Thr	His	Val	Glu	Ser	Pro	Trp	Arg	Leu	Ile
		515					520					525			
Arg	Pro	Ser	Val	Ile	Ser	Ile	Ile	Gly	Leu	Tyr	Lys	Glu	Lys	Gly	Lys
	530					535					540				
Gly	Leu	Gly	Phe	Ser	Ile	Ala	Gly	Gly	Arg	Asp	Cys	Ile	Arg	Gly	Gln
545					550					555					560
Met	Gly	Ile	Phe	Val	Lys	Thr	Ile	Phe	Pro	Asn	Gly	Ser	Ala	Ala	Glu
				565					570					575	
Asp	Gly	Arg	Leu	Lys	Glu	Gly	Asp	Glu	Ile	Leu	Asp	Val	Asn	Gly	Ile
			580					585					590		
Pro	Ile	Lys	Gly	Leu	Thr	Phe	Gln	Glu	Ala	Ile	His	Thr	Phe	Lys	Gln

595					600					605						
Ile	Arg	Ser	Gly	Leu	Phe	Val	Leu	Thr	Val	Arg	Thr	Lys	Leu	Leu	Ser	
610					615					620						
Pro	Ser	Leu	Thr	Pro	Cys	Ser	Thr	Pro	Thr	His	Met	Ser	Arg	Ser	Ser	
625					630					635					640	
Ser	Pro	Ser	Phe	Asn	Thr	Asn	Ser	Gly	Gly	Thr	Pro	Ala	Gly	Gly	Gly	
645					650					655						
Gln	Glu	Glu	Gly	Gly	Ser	Ser	Ser	Leu	Gly	Arg	Lys	Ala	Pro	Gly	Pro	
660					665					670						
Lys	Asp	Arg	Ile	Val	Met	Glu	Val	Thr	Leu	Asn	Lys	Glu	Pro	Arg	Val	
675					680					685						
Gly	Leu	Gly	Ile	Gly	Ala	Cys	Cys	Leu	Ala	Leu	Glu	Asn	Ser	Pro	Pro	
690					695					700						
Gly	Ile	Tyr	Ile	His	Ser	Leu	Ala	Pro	Gly	Ser	Val	Ala	Lys	Met	Glu	
705					710					715					720	
Ser	Asn	Leu	Ser	Arg	Gly	Asp	Gln	Ile	Leu	Glu	Val	Asn	Ser	Val	Asn	
725					730					735						
Val	Arg	His	Ala	Ala	Leu	Ser	Lys	Val	His	Ala	Ile	Leu	Ser	Lys	Cys	
740					745					750						
Pro	Pro	Gly	Pro	Val	Arg	Leu	Val	Ile	Gly	Arg	His	Pro	Asn	Pro	Lys	
755					760					765						
Val	Ser	Glu	Gln	Glu	Met	Asp	Glu	Val	Ile	Ala	Arg	Ser	Thr	Tyr	Gln	
770					775					780						
Glu	Ser	Arg	Glu	Ala	Asn	Ser	Ser	Pro	Gly	Leu	Gly	Thr	Pro	Leu	Lys	
785					790					795					800	
Ser	Pro	Ser	Leu	Ala	Lys	Lys	Asp	Ser	Leu	Leu	Ser	Glu	Ser	Glu	Leu	
805					810					815						
Ser	Gln	Tyr	Phe	Val	His	Asp	Gly	Gln	Gly	Ser	Leu	Ser	Asp	Phe	Val	
820					825					830						
Val	Ala	Gly	Ser	Glu	Asp	Glu	Asp	His	Pro	Gly	Ser	Gly	Tyr	Glu	Thr	
835					840					845						
Ser	Glu	Asp	Gly	Ser	Leu	Leu	Pro	Val	Pro	Ser	Ala	His	Lys	Ala	Arg	
850					855					860						
Ala	Asn	Ser	Leu	Val	Thr	Leu	Gly	Ser	Gln	Arg	Thr	Ser	Gly	Leu	Leu	
865					870					875					880	
His	Lys	Gln	Val	Thr	Val	Ala	Arg	Gln	Ala	Ser	Leu	Pro	Gly	Ser	Pro	
885					890					895						
Gln	Val	Leu	Arg	Asn	Pro	Leu	Leu	Arg	Gln	Arg	Arg	Val	Arg	Cys	Tyr	

900					905					910						
Asp	Ser	Asn	Gly	Gly	Ser	Asp	Asp	Glu	Asp	Phe	Asp	Gly	Glu	Gly	Asp	
915					920					925						
Cys	Ile	Ser	Leu	Pro	Gly	Val	Leu	Pro	Gly	Pro	Gly	Lys	Pro	Leu	Val	
930					935					940						
Glu	Asp	Asp	Thr	Arg	Pro	Ala	Leu	Thr	Thr	Ser	Ser	Lys	Ser	Ile	Asp	
945					950					955					960	
Val	Asn	Lys	Gln	Glu	Glu	Arg	Leu	Gln	Lys	Pro	Leu	Val	Ser	Lys	Ala	
965					970					975						
Cys	Ser	Val	Pro	Leu	Leu	Gly	Ser	Ser	Leu	Asp	Ser	Glu	His	Ser	Ile	
980					985					990						
Leu	Asn	Gly	Ala	Gly	Gly	Thr	Pro	Pro	Lys	Val	Ala	Ser	Leu	Pro	Gly	
995					1000					1005						
Ser	Gly	Glu	Thr	Pro	Lys	Asn	Gly	Pro	Arg	Gly	Ser	Gly	Arg	Lys	Glu	
1010					1015					1020						
Met	Ser	Gly	Ser	Arg	Ser	Ser	Pro	Lys	Leu	Glu	Tyr	Arg	Val	Pro	Thr	
1025					1030					1035					1040	
Asp	Thr	Gln	Ser	Pro	Arg	Ser	Pro	Glu	Asn	His	Thr	Ser	Pro	Pro	Gln	
1045					1050					1055						
Lys	Ser	Glu	Asn	Leu	Val	Ser	Arg	His	Lys	Pro	Val	Ala	Arg	Ile	Ser	
1060					1065					1070						
Pro	His	Tyr	Lys	Arg	Ser	Asp	Ala	Glu	Glu	Ala	Pro	Gly	Gly	Thr	Ala	
1075					1080					1085						
Asn	Gly	Pro	Cys	Ala	Gln	Asp	Leu	Lys	Val	Gln	Ala	Ser	Pro	Val	Lys	
1090					1095					1100						
Asp	Pro	Val	Thr	Ser	Arg	Gln	Pro	Gly	Gly	Thr	Ala	Glu	Lys	Glu	Leu	
1105					1110					1115					1120	
Arg	Gly	Asn	Pro	Thr	Pro	Gly	Asp	Ser	Ser	Val	Pro	Thr	Asn	Cys	Gly	
1125					1130					1135						
Pro	Ala	Ser	Thr	Pro	Cys	His	Pro	Asn	Ile	Gly	Leu	Pro	Thr	Glu	Asn	
1140					1145					1150						
Pro	Gln	Gly	Ala	Ala	Pro	Glu	Cys	Gly	Pro	His	Pro	Gly	Thr	Gly	Trp	
1155					1160					1165						
Asp	Gly	Ser	Ser	Glu	His	Leu	Cys	Ser	Pro	Gly	Lys	Ser	Arg	Glu	Val	
1170					1175					1180						
His	Pro	Asp	Ser	Ser	Glu	Thr	Pro	Thr	Val	Ala	Glu	Gln	Val	His	Gln	
1185					1190					1195					1200	
Pro	Glu	Ser	Leu	Ser	Gln	Pro	Val	Ser	Pro	Arg	Thr	Ser	Glu	Pro	Glu	

1205					1210					1215						
Ser	Gln	Gly	Ile	Ser	Lys	Met	Lys	Pro	Pro	Ser	Gln	Arg	Cys	Val	Ser	
1220					1225					1230						
Pro	Arg	Glu	Lys	Ala	Ser	Thr	Pro	Pro	Asp	Ser	Ser	Arg	Ala	Trp	Ala	
1235					1240					1245						
Ala	Pro	Gly	Asp	Ser	Ser	Pro	Ser	Thr	Arg	Arg	Ile	Ala	Val	Pro	Met	
1250					1255					1260						
Ser	Thr	Gly	Ala	Ala	Pro	Ala	Thr	Ala	Ile	Pro	Gln	Ala	Ser	Leu	Val	
1265					1270					1275					1280	
Ser	Gln	Glu	Arg	Ser	Arg	Gly	Leu	Ser	Gly	Pro	Ser	Lys	Gly	Leu	Gly	
1285					1290					1295						
Thr	Lys	Glu	Leu	Cys	Ile	Pro	Lys	Ser	Leu	Lys	Asp	Gly	Ala	Leu	Leu	
1300					1305					1310						
Glu	Asp	Thr	Ala	Pro	Ala	Ser	Gly	Lys	Met	Ser	His	Ala	Ser	Ser	Pro	
1315					1320					1325						
Ser	Gly	Pro	Val	Ala	Thr	Glu	Arg	Thr	Leu	Ser	Gly	Ser	Pro	Glu	Asn	
1330					1335					1340						
Pro	Val	Thr	Asp	Ile	Asp	Asn	Phe	Ile	Glu	Glu	Ala	Ser	Glu	Ala	Arg	
1345					1350					1355					1360	
Leu	Ser	Gln	Ser	Pro	Gln	Lys	Ala	Asp	Cys	Arg	Ala	His	Gly	Asp	Thr	
1365					1370					1375						
Phe	Glu	Ser	Gln	Pro	Pro	Gly	Gly	Ala	Gly	Ser	Ser	Ser	Ser	His	His	
1380					1385					1390						
Ala	Gln	Met	Val	Arg	Ser	Asp	Gln	Thr	Ser	Ser	Pro	Arg	Lys	Thr	Gly	
1395					1400					1405						
Gly	Thr	Gly	Ser	Pro	Pro	Pro	Gln	Gln	Trp	Ala	Leu	Gln	Pro	Ser	Val	
1410					1415					1420						
Leu	Asp	Ser	Ile	His	Pro	Asp	Lys	His	Leu	Ala	Val	Asn	Lys	Thr	Phe	
1425					1430					1435					1440	
Leu	Asn	Asn	Tyr	Ser	Arg	Asn	Phe	Ser	Asn	Phe	His	Glu	Asp	Ser	Ile	
1445					1450					1455						
Ser	Leu	Ser	Gly	Pro	Gly	Gly	Ser	Ser	Glu	Pro	Ser	Pro	Ser	Ser	Met	
1460					1465					1470						
Tyr	Gly	Asn	Ala	Glu	Asp	Ser	Ser	Ser	Asp	Pro	Glu	Ser	Leu	Ala	Glu	
1475					1480					1485						
Asp	Pro	Gly	Ala	Ala	Ala	Arg	Asn	Asn	Trp	Ser	Pro	Pro	Leu	Ser	Pro	
1490					1495					1500						
Glu	Ser	Ser	Pro	Lys	Glu	Gly	Ser	Ser	Glu	Ser	Glu	Asp	Glu	Arg	Ile	

1505	1510	1515	1520
Glu Ile Cys Ser Thr Asp Gly Cys Pro Gly Thr Pro Val Thr Ala Pro			
1525		1530	1535
Pro Pro Thr Gln Val Ala Leu Cys Pro Val Leu Pro Val Gln Gln Arg			
1540	1545		1550
Ala Val Cys Lys Pro Val Gly Asp Ile Cys Glu Arg Ala Cys Phe Val			
1555	1560		1565
Pro Gly Ala Ser Arg Thr Ser Ile Pro Asp Ser Ser Gln Pro Phe Ser			
1570	1575	1580	
Phe Leu Asp Val Ser Ser Glu Glu Pro Glu Thr Trp Ala Ser Ile Asn			
1585	1590	1595	1600
Ala Ser Gln Asn His Met Pro Val Cys Thr Glu Gly Ile Met Asp Val			
1605	1610		1615
Thr Ser Thr Ser Ser Asn Met Gly Asp Ser Gln Ser Ser Gln Met Thr			
1620	1625		1630
Arg His Cys Arg Asn Ala Pro Phe Val Leu Gly Asn Pro Asp Met Val			
1635	1640		1645
Asn Asp Leu Gly Arg Asp Leu Leu Asp Glu Gly Ala Pro Lys Glu Gly			
1650	1655	1660	
Ala Ala Ala Ala Ser Val Met Arg Ser Val Phe Ala Leu Gly Ala Glu			
1665	1670	1675	1680
Gly Pro Lys Asn Gly Glu Ala Val Leu Ala Asp Leu His Ile Ala Glu			
1685	1690		1695
Arg Gly Asn Leu Glu Asp Leu Leu Gln Lys Pro Lys Thr Ile Ser Arg			
1700	1705		1710
Arg Pro Ile Leu Thr Trp Phe Lys Glu Ile Asn Lys Asp Ser Gln Gly			
1715	1720		1725
Ser His Leu Arg Ser Thr Ser Glu Lys Glu Gln Ser Ser Met Leu Ala			
1730	1735	1740	
Leu Gly Pro Gly Ser Lys Ala Asn Met Val Asn Thr Gly His Arg Lys			
1745	1750	1755	1760
Gly Val Thr Val Pro Lys Ser Pro Pro Ser Arg Gln Lys Ser Gln Glu			
1765	1770		1775
Asn Lys Asp Leu Pro Pro Lys Ser Pro Val Glu Thr Leu Gly Asn Cys			
1780	1785		1790
Gln Lys Pro Lys Cys Ser Pro Lys Leu Lys Arg Leu Asn Ser Lys Gly			
1795	1800		1805
Lys Ala Ser Pro Glu Val Pro Val Ala Ile Ser Thr Lys Gly Ser Arg			

1810	1815	1820
Asn Asp His Arg Lys Thr Leu Pro Ser Pro Gln Ala Ser His Lys Met		
1825	1830	1835 1840
Phe Ser Lys Ala Val Ser His Arg Leu His Ile Ala Asp Gln Glu Glu		
	1845	1850 1855
Pro Lys Asn Thr Ala Gly Asp Thr Pro Lys Pro Pro Gln Cys Val Pro		
	1860	1865 1870
Glu Ser Lys Pro Pro Gln Ala Ala Leu Gly Ser Leu Arg Thr Ser Ala		
	1875	1880 1885
Ser Asp Thr Ser Ile Arg Thr Phe Thr Ser Pro Leu Thr Ser Pro Lys		
	1890	1895 1900
Leu Leu Pro Glu Gln Gly Ala Asn Ser Arg Phe His Met Ala Val Tyr		
	1905	1910 1915 1920
Leu Glu Ser Asp Thr Ser Cys Pro Thr Thr Ser Arg Ser Pro Arg Ser		
	1925	1930 1935
Gly Pro Glu Gly Lys Ala Pro His Ala Asn Ser Gly Ser Ala Ser Pro		
	1940	1945 1950
Pro Ala Ser Arg Ala Ser Leu Ala Leu Ala Gly Ile Arg Gln Ser Lys		
	1955	1960 1965
Gln Phe Thr Pro Gly Arg Ala Asp Leu Leu Val Ser Glu Ala Thr Gln		
	1970	1975 1980
Pro Gln Gly Ile Cys Glu Lys Gly Ala Glu Lys Lys Val Ser Asp Pro		
	1985	1990 1995 2000
Pro Gln Arg Thr Asn Gln Leu Lys Ile Val Glu Ile Ser Ser Glu Arg		
	2005	2010 2015
Val Pro Lys Asn Ala Cys Gly Asp Arg Pro Pro Glu Ser Asp Arg Lys		
	2020	2025 2030
Gly Gly Phe Leu Thr Gln Asn Asn Cys Gln Glu Lys Ser Ala Ile Arg		
	2035	2040 2045
Leu Arg Gln Ser Glu Glu Ser Ser Pro Glu His Thr Pro Phe Pro Pro		
	2050	2055 2060
Ser Gln Ala Ser Gln Val Glu Arg Glu Ile Arg Trp Ser Phe Ser Met		
	2065	2070 2075 2080
Ala Lys Pro Ala Thr Ser Ser Ser Ser Ser Leu Gln Leu Pro Ala Lys		
	2085	2090 2095
Leu Pro Glu Ser Phe Gln Gly Lys Ser Ser Gln Met Pro Ala Ser Val		
	2100	2105 2110
Gly Val Pro Lys Asn Gly Val Pro Ile Gly Leu Ala Gly Glu Glu Ser		

2115	2120	2125
Pro Tyr Phe Thr Pro Arg Pro Ala Thr Arg Thr Tyr Ser Met Pro Ala 2130	2135	2140
Gln Phe Ser Ser His Phe Gly Arg Glu Gly Pro Ser Pro His Ser Pro 2145	2150	2155 2160
Ser His Ser Pro Gln Asp Pro Gln Val Pro Ala Met Gly Gly Lys Leu 2165	2170	2175
Ser Glu Lys Thr Ala Lys Gly Val Thr Asn Gly Gln Gly Val Tyr Ser 2180	2185	2190
Val Lys Pro Leu Leu Glu Thr Ser Lys Asn Leu Ser Pro Val Asp Gly 2195	2200	2205
Arg Asp Val Ser Ala Asp Pro Glu Thr Ser Cys Leu Ile Pro Asp Lys 2210	2215	2220
Val Lys Val Thr Arg Arg Gln Tyr Cys Cys Glu Gln Ser Trp Pro His 2225	2230	2235 2240
Glu Ser Thr Ser Phe Phe Ser Val Lys Gln Arg Ile Lys Ser Phe Glu 2245	2250	2255
Asn Leu Ala Asn Ser Asp Arg Pro Thr Ala Lys Cys Ala Thr Ser Pro 2260	2265	2270
Phe Leu Ser Val Ser Ser Lys Pro Pro Ile Asn Arg Arg Ser Ser Gly 2275	2280	2285
Ser Ile Pro Ser Gly Ser Pro Ser Asp Met Thr Ser Arg Ser Leu Arg 2290	2295	2300
Arg Ser Leu Ser Ser Cys Ser Glu Ser Gln Ser Glu Ala Ser Ser Leu 2305	2310	2315 2320
Leu Pro Gln Met Thr Lys Ser Pro Ser Ser Met Thr Leu Thr Val Ser 2325	2330	2335
Arg Gln Asn Pro Pro Asp Thr Ser Asn Lys Gly Pro Ser Pro Asp Pro 2340	2345	2350
Lys Lys Ser Leu Val Pro Val Gly Ile Pro Thr Ser Thr Val Ser Pro 2355	2360	2365
Ala Ser Pro Ser Lys Arg Asn Lys Ser Ser Val Arg His Ala Gln Pro 2370	2375	2380
Ser Pro Val Ser Arg Ser Lys Leu Gln Glu Arg Arg Thr Leu Ser Met 2385	2390	2395 2400
Pro Asp Leu Asp Lys Leu Cys Asn Gly Glu Asp Asp Ser Ala Ser Pro 2405	2410	2415
Gly Ala Val Leu Phe Lys Thr Gln Leu Glu Ile Thr Pro Arg Arg Ser		

2420	2425	2430
Lys Gly Ser Gln Ala Thr Ser Pro Ala Gly Ser Pro Ala Arg Gly His		
2435	2440	2445
Ala Asp Phe Asn Gly Ser Thr Phe Leu Ser Cys Pro Met Asn Gly Gly		
2450	2455	2460
Thr Arg Ala Tyr Thr Lys Gly Asn Ser Pro Pro Ala Ser Glu Pro Ala		
2465	2470	2475 2480
Ile Ala Thr Gly Ser Arg Glu Glu Gly Glu Ser Val Trp Ala Thr Pro		
2485	2490	2495
Ser Gly Lys Ser Trp Ser Val Ser Leu Asp Arg Leu Leu Ala Ser Val		
2500	2505	2510
Gly Asn Gln Gln Arg Leu Gln Gly Ile Leu Ser Leu Val Gly Ser Lys		
2515	2520	2525
Ser Pro Ile Leu Thr Leu Ile Gln Glu Ala Lys Ala Gln Ser Glu Thr		
2530	2535	2540
Lys Glu Asp Ile Cys Phe Ile Val Leu Asn Lys Lys Glu Gly Ser Gly		
2545	2550	2555 2560
Leu Gly Phe Ser Val Ala Gly Gly Ala Asp Val Glu Pro Lys Ser Val		
2565	2570	2575
Met Val His Arg Val Phe Ser Gln Gly Val Ala Ser Gln Glu Gly Thr		
2580	2585	2590
Val Ser Arg Gly Asp Phe Leu Leu Ser Val Asn Gly Thr Ser Leu Ala		
2595	2600	2605
Gly Leu Ala His Ser Glu Val Thr Lys Val Leu His Gln Ala Glu Leu		
2610	2615	2620
His Lys His Ala Leu Met Ile Ile Lys Lys Gly Asn Asp Gln Pro Gly		
2625	2630	2635 2640
Pro Ser Phe Lys Gln Glu Pro Pro Ser Ala Asn Gly Lys Gly Pro Phe		
2645	2650	2655
Pro Arg Arg Thr Leu Pro Leu Glu Pro Gly Ala Gly Arg Asn Gly Ala		
2660	2665	2670
Ala His Asp Ala Leu Cys Val Glu Val Leu Lys Thr Ser Ala Gly Leu		
2675	2680	2685
Gly Leu Ser Leu Asp Gly Gly Lys Ser Ser Val Ser Gly Glu Gly Pro		
2690	2695	2700
Leu Val Ile Lys Arg Val Tyr Lys Gly Gly Ala Ala Glu Arg Ala Gly		
2705	2710	2715 2720
Thr Ile Glu Ala Gly Asp Glu Ile Leu Ala Ile Asn Gly Lys Pro Leu		

2725	2730	2735
Val Gly Leu Val His Phe Asp Ala Trp Asn Ile Met Lys Ser Val Pro		
2740	2745	2750
Glu Gly Pro Val Gln Leu Val Ile Arg Lys His Arg Asp Ser		
2755	2760	2765
<210> 63		
<211> 2641		
<212> PRT		
<213> Homo sapiens		
<400> 63		
Met Pro Ile Thr Gln Asp Asn Ala Val Leu His Leu Pro Leu Leu Tyr		
1	5	10 15
Gln Trp Leu Gln Asn Ser Leu Gln Glu Gly Gly Asp Gly Pro Glu Gln		
20	25	30
Arg Leu Cys Gln Ala Ala Ile Gln Lys Leu Gln Glu Tyr Ile Gln Leu		
35	40	45
Asn Phe Ala Val Asp Glu Ser Thr Val Pro Pro Asp His Ser Pro Pro		
50	55	60
Glu Met Glu Ile Cys Thr Val Tyr Leu Thr Lys Glu Leu Gly Asp Thr		
65	70	75 80
Glu Thr Val Gly Leu Ser Phe Gly Asn Ile Pro Val Phe Gly Asp Tyr		
85	90	95
Gly Glu Lys Arg Arg Gly Gly Lys Lys Arg Lys Thr His Gln Gly Pro		
100	105	110
Val Leu Asp Val Gly Cys Ile Trp Val Thr Glu Leu Arg Lys Asn Ser		
115	120	125
Pro Ala Gly Lys Ser Gly Lys Val Arg Leu Arg Asp Glu Ile Leu Ser		
130	135	140
Leu Asn Gly Gln Leu Met Val Gly Val Asp Val Ser Gly Ala Ser Tyr		
145	150	155 160
Leu Ala Glu Gln Cys Trp Asn Gly Gly Phe Ile Tyr Leu Ile Met Leu		
165	170	175
Arg Arg Phe Lys His Lys Ala His Ser Thr Tyr Asn Gly Asn Ser Ser		
180	185	190
Asn Ser Ser Glu Pro Gly Glu Thr Pro Thr Leu Glu Leu Gly Asp Arg		
195	200	205
Thr Ala Lys Lys Gly Lys Arg Thr Arg Lys Phe Gly Val Ile Ser Arg		
210	215	220

Pro Pro Ala Asn Lys Ala Pro Glu Glu Ser Lys Gly Ser Ala Gly Cys
 225 230 235 240
 Glu Val Ser Ser Asp Pro Ser Thr Glu Leu Glu Asn Gly Ala Asp Pro
 245 250 255
 Glu Leu Gly Asn Gly His Val Phe Gln Leu Glu Asn Gly Pro Asp Ser
 260 265 270
 Leu Lys Glu Val Ala Gly Pro His Leu Glu Arg Ser Glu Val Asp Arg
 275 280 285
 Gly Thr Glu His Arg Ile Pro Lys Thr Asp Ala Pro Leu Thr Thr Ser
 290 295 300
 Asn Asp Lys Arg Arg Phe Ser Lys Gly Gly Lys Thr Asp Phe Gln Ser
 305 310 315 320
 Ser Asp Cys Leu Ala Arg Gln Glu Glu Val Gly Arg Ile Trp Lys Met
 325 330 335
 Glu Leu Leu Lys Glu Ser Asp Gly Leu Gly Ile Gln Val Ser Gly Gly
 340 345 350
 Arg Gly Ser Lys Arg Ser Pro His Ala Ile Val Val Thr Gln Val Lys
 355 360 365
 Glu Gly Gly Ala Ala His Arg Glu Tyr His Ile Val Lys Lys Ser Thr
 370 375 380
 Arg Ser Leu Ser Thr Thr Gln Val Glu Ser Pro Trp Arg Leu Ile Arg
 385 390 395 400
 Pro Ser Val Ile Ser Ile Ile Gly Leu Tyr Lys Glu Lys Gly Lys Gly
 405 410 415
 Leu Gly Phe Ser Ile Ala Gly Gly Arg Asp Cys Ile Arg Gly Gln Met
 420 425 430
 Gly Ile Phe Val Lys Thr Ile Phe Pro Asn Gly Ser Ala Ala Glu Asp
 435 440 445
 Gly Arg Leu Lys Glu Gly Asp Glu Ile Leu Asp Val Asn Gly Ile Pro
 450 455 460
 Ile Lys Gly Leu Thr Phe Gln Glu Ala Ile His Thr Phe Lys Gln Ile
 465 470 475 480
 Arg Ser Gly Leu Phe Val Leu Thr Val Arg Thr Lys Leu Val Ser Pro
 485 490 495
 Ser Leu Thr Pro Cys Ser Thr Pro Thr His Met Ser Arg Ser Ala Ser
 500 505 510
 Pro Asn Phe Asn Thr Ser Gly Gly Ala Ser Ala Gly Gly Ser Asp Glu
 515 520 525

Gly	Ser	Ser	Ser	Ser	Leu	Gly	Arg	Lys	Thr	Pro	Gly	Pro	Lys	Asp	Arg		
530						535					540						
Ile	Val	Met	Glu	Val	Thr	Leu	Asn	Lys	Glu	Pro	Arg	Val	Gly	Leu	Gly		
545					550					555					560		
Ile	Gly	Ala	Cys	Cys	Leu	Ala	Leu	Glu	Asn	Ser	Pro	Pro	Gly	Ile	Tyr		
			565						570					575			
Ile	His	Ser	Leu	Ala	Pro	Gly	Ser	Val	Ala	Lys	Met	Glu	Ser	Asn	Leu		
			580					585					590				
Ser	Arg	Gly	Asp	Gln	Ile	Leu	Glu	Val	Asn	Ser	Val	Asn	Val	Arg	His		
		595					600					605					
Ala	Ala	Leu	Ser	Lys	Val	His	Ala	Ile	Leu	Ser	Lys	Cys	Pro	Val	Ser		
610						615					620						
Glu	Gln	Glu	Met	Asp	Glu	Val	Ile	Ala	Arg	Ser	Thr	Tyr	Gln	Glu	Ser		
625					630					635					640		
Lys	Glu	Ala	Asn	Ser	Ser	Pro	Gly	Leu	Gly	Thr	Pro	Leu	Lys	Ser	Pro		
			645						650					655			
Ser	Leu	Ala	Lys	Lys	Asp	Ser	Leu	Ile	Ser	Glu	Ser	Glu	Leu	Ser	Gln		
			660					665					670				
Tyr	Phe	Ala	His	Asp	Val	Pro	Gly	Pro	Leu	Ser	Asp	Phe	Met	Val	Ala		
	675						680					685					
Gly	Ser	Glu	Asp	Glu	Asp	His	Pro	Gly	Ser	Gly	Cys	Ser	Thr	Ser	Glu		
690						695					700						
Glu	Gly	Ser	Leu	Pro	Pro	Ser	Thr	Ser	Ser	Glu	Pro	Gly	Lys	Pro	Arg		
705					710					715					720		
Ala	Asn	Ser	Leu	Val	Thr	Leu	Gly	Ser	His	Arg	Ala	Ser	Gly	Leu	Phe		
				725					730					735			
His	Lys	Gln	Val	Thr	Val	Ala	Arg	Gln	Ala	Ser	Leu	Pro	Gly	Ser	Pro		
			740					745					750				
Gln	Ala	Leu	Arg	Asn	Pro	Leu	Leu	Arg	Gln	Arg	Lys	Val	Gly	Cys	Tyr		
		755					760					765					
Asp	Ala	Asn	Asp	Ala	Ser	Asp	Glu	Glu	Glu	Phe	Asp	Arg	Glu	Gly	Asp		
770						775					780						
Cys	Ile	Ser	Leu	Pro	Gly	Ala	Leu	Pro	Gly	Pro	Ile	Arg	Pro	Leu	Ser		
785					790					795					800		
Glu	Asp	Asp	Pro	Arg	Arg	Val	Ser	Ile	Ser	Ser	Ser	Lys	Gly	Met	Asp		
			805						810					815			
Val	His	Asn	Gln	Glu	Glu	Arg	Pro	Arg	Lys	Thr	Leu	Val	Ser	Lys	Ala		
			820					825						830			

Ile	Ser	Ala	Pro	Leu	Leu	Gly	Ser	Ser	Val	Asp	Leu	Glu	Glu	Ser	Ile	835	840	845
Pro	Glu	Gly	Met	Val	Asp	Ala	Ala	Ser	Tyr	Ala	Ala	Asn	Leu	Thr	Asp	850	855	860
Ser	Ala	Glu	Ala	Pro	Lys	Gly	Ser	Pro	Gly	Ser	Trp	Trp	Lys	Lys	Glu	865	870	875
Leu	Ser	Gly	Ser	Ser	Ser	Ala	Pro	Lys	Leu	Glu	Tyr	Thr	Val	Arg	Thr	885	890	895
Asp	Thr	Gln	Ser	Pro	Thr	Asn	Thr	Gly	Ser	Pro	Ser	Ser	Pro	Gln	Gln	900	905	910
Lys	Ser	Glu	Gly	Leu	Gly	Ser	Arg	His	Arg	Pro	Val	Ala	Arg	Val	Ser	915	920	925
Pro	His	Cys	Lys	Arg	Ser	Glu	Ala	Glu	Ala	Lys	Pro	Ser	Gly	Ser	Gln	930	935	940
Thr	Val	Asn	Leu	Thr	Gly	Arg	Ala	Asn	Asp	Pro	Cys	Asp	Leu	Asp	Ser	945	950	955
Arg	Val	Gln	Ala	Thr	Ser	Val	Lys	Val	Thr	Val	Ala	Gly	Phe	Gln	Pro	965	970	975
Gly	Gly	Ala	Val	Glu	Lys	Leu	Cys	Gln	Glu	Ser	Leu	Gly	Lys	Leu	Thr	980	985	990
Thr	Gly	Asp	Ala	Cys	Val	Ser	Thr	Ser	Cys	Glu	Leu	Ala	Ser	Ala	Leu	995	1000	1005
Ser	His	Leu	Asp	Ala	Ser	His	Leu	Thr	Glu	Asn	Leu	Pro	Lys	Ala	Ala	1010	1015	1020
Ser	Glu	Leu	Gly	Gln	Gln	Pro	Met	Thr	Ser	Ser	Asp	Leu	Ile	Ser	Ser	1025	1030	1035
Pro	Gly	Lys	Lys	Gly	Ala	Ala	His	Pro	Asp	Pro	Ser	Lys	Thr	Ser	Val	1045	1050	1055
Asp	Thr	Gly	Gln	Val	Ser	Arg	Pro	Glu	Asn	Pro	Ser	Gln	Pro	Ala	Ser	1060	1065	1070
Pro	Arg	Val	Thr	Lys	Cys	Lys	Ala	Arg	Ser	Pro	Val	Arg	Leu	Pro	His	1075	1080	1085
Glu	Gly	Ser	Pro	Ser	Pro	Gly	Glu	Lys	Ala	Ala	Ala	Pro	Pro	Asp	Tyr	1090	1095	1100
Ser	Lys	Thr	Arg	Ser	Ala	Ser	Glu	Thr	Ser	Thr	Pro	His	Asn	Thr	Arg	1105	1110	1115
Arg	Val	Ala	Ala	Leu	Arg	Gly	Ala	Gly	Pro	Gly	Ala	Glu	Gly	Met	Thr	1125	1130	1135

Pro Ala Gly Ala Val Leu Pro Gly Asp Pro Leu Thr Ser Gln Glu Gln
 1140 1145 1150
 Arg Gln Gly Ala Pro Gly Asn His Ser Lys Ala Leu Glu Met Thr Gly
 1155 1160 1165
 Ile His Ala Pro Glu Ser Ser Gln Glu Pro Ser Leu Leu Glu Gly Ala
 1170 1175 1180
 Asp Ser Val Ser Ser Arg Ala Pro Gln Ala Ser Leu Ser Met Leu Pro
 1185 1190 1195 1200
 Ser Thr Asp Asn Thr Lys Glu Ala Cys Gly His Val Ser Gly His Cys
 1205 1210 1215
 Cys Pro Gly Gly Ser Arg Glu Ser Pro Val Thr Asp Ile Asp Ser Phe
 1220 1225 1230
 Ile Lys Glu Leu Asp Ala Ser Ala Ala Arg Ser Pro Ser Ser Gln Thr
 1235 1240 1245
 Gly Asp Ser Gly Ser Gln Glu Gly Ser Ala Gln Gly His Pro Pro Ala
 1250 1255 1260
 Gly Ala Gly Gly Gly Ser Ser Cys Arg Ala Glu Pro Val Pro Gly Gly
 1265 1270 1275 1280
 Gln Thr Ser Ser Pro Arg Arg Ala Trp Ala Ala Gly Ala Pro Ala Tyr
 1285 1290 1295
 Pro Gln Trp Ala Ser Gln Pro Ser Val Leu Asp Ser Ile Asn Pro Asp
 1300 1305 1310
 Lys His Phe Thr Val Asn Lys Asn Phe Leu Ser Asn Tyr Ser Arg Asn
 1315 1320 1325
 Phe Ser Ser Phe His Glu Asp Ser Thr Ser Leu Ser Gly Leu Gly Asp
 1330 1335 1340
 Ser Thr Glu Pro Ser Leu Ser Ser Met Tyr Gly Asp Ala Glu Asp Ser
 1345 1350 1355 1360
 Ser Ser Asp Pro Glu Ser Leu Thr Glu Ala Pro Arg Ala Ser Ala Arg
 1365 1370 1375
 Asp Gly Trp Ser Pro Pro Arg Ser Arg Val Ser Leu His Lys Glu Asp
 1380 1385 1390
 Pro Ser Glu Ser Glu Glu Glu Gln Ile Glu Ile Cys Ser Thr Arg Gly
 1395 1400 1405
 Cys Pro Asn Pro Pro Ser Ser Pro Ala His Leu Pro Thr Gln Ala Ala
 1410 1415 1420
 Ile Cys Pro Ala Ser Ala Lys Val Leu Ser Leu Lys Tyr Ser Thr Pro
 1425 1430 1435 1440

Arg Glu Ser Val Ala Ser Pro Arg Glu Lys Ala Ala Cys Leu Pro Gly
 1445 1450 1455
 Ser Tyr Thr Ser Gly Pro Asp Ser Ser Gln Pro Ser Ser Leu Leu Glu
 1460 1465 1470
 Met Ser Ser Gln Glu His Glu Thr His Ala Asp Ile Ser Thr Ser Gln
 1475 1480 1485
 Asn His Arg Pro Ser Cys Ala Glu Glu Thr Thr Glu Val Thr Ser Ala
 1490 1495 1500
 Ser Ser Ala Met Glu Asn Ser Pro Leu Ser Lys Val Ala Arg His Phe
 1505 1510 1515 1520
 His Ser Pro Pro Ile Ile Leu Ser Ser Pro Asn Met Val Asn Gly Leu
 1525 1530 1535
 Glu His Asp Leu Leu Asp Asp Glu Thr Leu Asn Gln Tyr Glu Thr Ser
 1540 1545 1550
 Ile Asn Ala Ala Ala Ser Leu Ser Ser Phe Ser Val Asp Val Pro Lys
 1555 1560 1565
 Asn Gly Glu Ser Val Leu Glu Asn Leu His Ile Ser Glu Ser Gln Asp
 1570 1575 1580
 Leu Asp Asp Leu Leu Gln Lys Pro Lys Met Ile Ala Arg Arg Pro Ile
 1585 1590 1595 1600
 Met Ala Trp Phe Lys Glu Ile Asn Lys His Asn Gln Gly Thr His Leu
 1605 1610 1615
 Arg Ser Lys Thr Glu Lys Glu Gln Pro Leu Met Pro Ala Arg Ser Pro
 1620 1625 1630
 Asp Ser Lys Ile Gln Met Val Ser Ser Ser Gln Lys Lys Gly Val Thr
 1635 1640 1645
 Val Pro His Ser Pro Pro Gln Pro Lys Thr Asn Leu Glu Asn Lys Asp
 1650 1655 1660
 Leu Ser Lys Lys Ser Pro Ala Glu Met Leu Leu Thr Asn Gly Gln Lys
 1665 1670 1675 1680
 Ala Lys Cys Gly Pro Lys Leu Lys Arg Leu Ser Leu Lys Gly Lys Ala
 1685 1690 1695
 Lys Val Asn Ser Glu Ala Pro Ala Ala Asn Ala Val Lys Ala Gly Gly
 1700 1705 1710
 Thr Asp His Arg Lys Pro Leu Ile Ser Pro Gln Thr Ser His Lys Thr
 1715 1720 1725
 Leu Ser Lys Ala Val Ser Gln Arg Leu His Val Ala Asp His Glu Asp
 1730 1735 1740

Pro Asp Arg Asn Thr Thr Ala Ala Pro Arg Ser Pro Gln Cys Val Leu
 1745 1750 1755 1760

Glu Ser Lys Pro Pro Leu Ala Thr Ser Gly Pro Leu Lys Pro Ser Val
 1765 1770 1775

Ser Asp Thr Ser Ile Arg Thr Phe Val Ser Pro Leu Thr Ser Pro Lys
 1780 1785 1790

Pro Val Pro Glu Gln Gly Met Trp Ser Arg Phe His Met Ala Val Leu
 1795 1800 1805

Ser Glu Pro Asp Arg Gly Cys Pro Thr Thr Pro Lys Ser Pro Lys Cys
 1810 1815 1820

Arg Ala Glu Gly Arg Ala Pro Arg Ala Asp Ser Gly Pro Val Ser Pro
 1825 1830 1835 1840

Ala Ala Ser Arg Asn Gly Met Ser Val Ala Gly Asn Arg Gln Ser Glu
 1845 1850 1855

Pro Arg Leu Ala Ser His Val Ala Ala Asp Thr Ala Gln Pro Arg Pro
 1860 1865 1870

Thr Gly Glu Lys Gly Gly Asn Ile Met Ala Ser Asp Arg Leu Glu Arg
 1875 1880 1885

Thr Asn Gln Leu Lys Ile Val Glu Ile Ser Ala Glu Ala Val Ser Glu
 1890 1895 1900

Thr Val Cys Gly Asn Lys Pro Ala Glu Ser Asp Arg Arg Gly Gly Cys
 1905 1910 1915 1920

Leu Ala Gln Gly Asn Cys Gln Glu Lys Ser Glu Ile Arg Leu Tyr Arg
 1925 1930 1935

Gln Val Ala Glu Ser Ser Thr Ser His Pro Ser Ser Leu Pro Ser His
 1940 1945 1950

Ala Ser Gln Ala Glu Gln Glu Met Ser Arg Ser Phe Ser Met Ala Lys
 1955 1960 1965

Leu Ala Ser Ser Ser Ser Ser Leu Gln Thr Ala Ile Arg Lys Ala Glu
 1970 1975 1980

Tyr Ser Gln Gly Lys Ser Ser Leu Met Ser Asp Ser Arg Gly Val Pro
 1985 1990 1995 2000

Arg Asn Ser Ile Pro Gly Gly Pro Ser Gly Glu Asp His Leu Tyr Phe
 2005 2010 2015

Thr Pro Arg Pro Ala Thr Arg Thr Tyr Ser Met Pro Ala Gln Phe Ser
 2020 2025 2030

Ser His Phe Gly Arg Glu Gly His Pro Pro His Ser Leu Gly Arg Ser
 2035 2040 2045

Arg Asp Ser Gln Val Pro Val Thr Ser Ser Val Val Pro Glu Ala Lys
 2050 2055 2060
 Ala Ser Arg Gly Gly Leu Pro Ser Leu Ala Asn Gly Gln Gly Ile Tyr
 2065 2070 2075 2080
 Ser Val Lys Pro Leu Leu Asp Thr Ser Arg Asn Leu Pro Ala Thr Asp
 2085 2090 2095
 Glu Gly Asp Ile Ile Ser Val Gln Glu Thr Ser Cys Leu Val Thr Asp
 2100 2105 2110
 Lys Ile Lys Val Thr Arg Arg His Tyr Cys Tyr Glu Gln Asn Trp Pro
 2115 2120 2125
 His Glu Ser Thr Ser Phe Phe Ser Val Lys Gln Arg Ile Lys Ser Phe
 2130 2135 2140
 Glu Asn Leu Ala Asn Ala Asp Arg Pro Val Ala Lys Ser Gly Ala Ser
 2145 2150 2155 2160
 Pro Phe Leu Ser Val Ser Ser Lys Pro Pro Ile Gly Arg Arg Ser Ser
 2165 2170 2175
 Gly Ser Ile Val Ser Gly Ser Leu Gly His Pro Gly Asp Ala Ala Ala
 2180 2185 2190
 Arg Leu Leu Arg Arg Ser Leu Ser Ser Cys Ser Glu Asn Gln Ser Glu
 2195 2200 2205
 Ala Gly Thr Leu Leu Pro Gln Met Ala Lys Ser Pro Ser Ile Met Thr
 2210 2215 2220
 Leu Thr Ile Ser Arg Gln Asn Pro Pro Glu Thr Ser Ser Lys Gly Ser
 2225 2230 2235 2240
 Asp Ser Glu Leu Lys Lys Ser Leu Gly Pro Leu Gly Ile Pro Thr Pro
 2245 2250 2255
 Thr Met Thr Leu Ala Ser Pro Val Lys Arg Asn Lys Ser Ser Val Arg
 2260 2265 2270
 His Thr Gln Pro Ser Pro Val Ser Arg Ser Lys Leu Gln Glu Leu Arg
 2275 2280 2285
 Ala Leu Ser Met Pro Asp Leu Asp Lys Leu Cys Ser Glu Asp Tyr Ser
 2290 2295 2300
 Ala Gly Pro Ser Ala Val Leu Phe Lys Thr Glu Leu Glu Ile Thr Pro
 2305 2310 2315 2320
 Arg Arg Ser Pro Gly Pro Pro Ala Gly Gly Val Ser Cys Pro Glu Lys
 2325 2330 2335
 Gly Gly Asn Arg Ala Cys Pro Gly Gly Ser Gly Pro Lys Thr Ser Ala
 2340 2345 2350

Ala Glu Thr Pro Ser Ser Ala Ser Asp Thr Gly Glu Ala Ala Gln Asp
 2355 2360 2365
 Leu Pro Phe Arg Arg Ser Trp Ser Val Lys Leu Asp Gln Leu Leu Val
 2370 2375 2380
 Ser Ala Gly Asp Gln Gln Arg Leu Gln Ser Val Leu Ser Ser Val Gly
 2385 2390 2395 2400
 Ser Lys Ser Thr Ile Leu Thr Leu Ile Gln Glu Ala Lys Ala Gln Ser
 2405 2410 2415
 Glu Asn Glu Glu Asp Val Cys Phe Ile Val Leu Asn Arg Lys Glu Gly
 2420 2425 2430
 Ser Gly Leu Gly Phe Ser Val Ala Gly Gly Thr Asp Val Glu Pro Lys
 2435 2440 2445
 Ser Ile Thr Val His Arg Val Phe Ser Gln Gly Ala Ala Ser Gln Glu
 2450 2455 2460
 Gly Thr Met Asn Arg Gly Asp Phe Leu Leu Ser Val Asn Gly Ala Ser
 2465 2470 2475 2480
 Leu Ala Gly Leu Ala His Gly Asn Val Leu Lys Val Leu His Gln Ala
 2485 2490 2495
 Gln Leu His Lys Asp Ala Leu Val Val Ile Lys Lys Gly Met Asp Gln
 2500 2505 2510
 Pro Arg Pro Ser Ala Arg Gln Glu Pro Pro Thr Ala Asn Gly Lys Gly
 2515 2520 2525
 Leu Leu Ser Arg Lys Thr Ile Pro Leu Glu Pro Gly Ile Gly Arg Ser
 2530 2535 2540
 Val Ala Val His Asp Ala Leu Cys Val Glu Val Leu Lys Thr Ser Ala
 2545 2550 2555 2560
 Gly Leu Gly Leu Ser Leu Asp Gly Gly Lys Ser Ser Val Thr Gly Asp
 2565 2570 2575
 Gly Pro Leu Val Ile Lys Arg Val Tyr Lys Gly Gly Ala Ala Glu Gln
 2580 2585 2590
 Ala Gly Ile Ile Glu Ala Gly Asp Glu Ile Leu Ala Ile Asn Gly Lys
 2595 2600 2605
 Pro Leu Val Gly Leu Met His Phe Asp Ala Trp Asn Ile Met Lys Ser
 2610 2615 2620
 Val Pro Glu Gly Pro Val Gln Leu Leu Ile Arg Lys His Arg Asn Ser
 2625 2630 2635 2640
 Ser

<210> 64
 <211> 364
 <212> PRT
 <213> Mus musculus

<400> 64

Met	Leu	Arg	Arg	Phe	Lys	Gln	Lys	Ala	His	Leu	Thr	Tyr	Asn	Gly	Asn
1				5					10					15	
Ser	Gly	Asn	Ser	Ser	Glu	Pro	Gly	Glu	Thr	Pro	Thr	Leu	Glu	Leu	Gly
		20						25					30		
Asp	Gln	Thr	Ser	Lys	Lys	Gly	Lys	Arg	Thr	Arg	Lys	Phe	Gly	Val	Ile
		35					40					45			
Ser	Arg	Pro	Ala	Ile	Ile	Lys	Ala	Pro	Glu	Asp	Ser	Lys	Ser	Asn	Ser
	50					55					60				
Gly	Cys	Asp	Thr	Ala	Asp	Asp	Pro	Ser	Ser	Glu	Leu	Glu	Asn	Gly	Thr
65					70					75					80
Asp	Ser	Glu	Leu	Gly	Asn	Gly	His	Ala	Phe	Glu	Leu	Glu	Asn	Gly	Pro
				85					90					95	
Asn	Ser	Leu	Lys	Asp	Val	Ala	Gly	Pro	His	Leu	Glu	Arg	Ser	Glu	Ala
		100						105					110		
Asp	Arg	Glu	Ala	Glu	Leu	Arg	Val	Pro	Lys	Thr	Glu	Ala	Pro	Leu	Ser
		115					120					125			
Asp	Ser	Asn	Asp	Lys	Arg	Arg	Phe	Ser	Lys	Thr	Gly	Lys	Thr	Asn	Phe
	130					135					140				
Gln	Ser	Ser	Asp	Ser	Leu	Ala	Arg	Glu	Glu	Val	Gly	Arg	Ile	Trp	Glu
145					150					155					160
Met	Glu	Leu	Leu	Lys	Glu	Ser	Asp	Gly	Leu	Gly	Ile	Gln	Val	Ser	Gly
				165					170					175	
Gly	Arg	Gly	Ser	Lys	Arg	Ser	Pro	His	Ala	Ile	Val	Val	Thr	Gln	Val
			180					185					190		
Lys	Glu	Gly	Gly	Ala	Ala	His	Arg	Asp	Gly	Arg	Leu	Ser	Leu	Gly	Asp
		195					200					205			
Glu	Leu	Leu	Val	Ile	Asn	Gly	His	Leu	Leu	Val	Gly	Leu	Ser	His	Glu
	210					215					220				
Glu	Ala	Val	Ala	Ile	Leu	Arg	Ser	Ala	Thr	Gly	Met	Val	Gln	Leu	Val
225					230					235					240
Val	Ala	Ser	Lys	Met	Leu	Gly	Ser	Glu	Glu	Ser	Gln	Asp	Val	Gly	Ser
				245					250					255	
Ser	Glu	Glu	Ser	Lys	Gly	Asn	Asn	Leu	Glu	Ser	Pro	Lys	Gln	Gly	Asn
			260					265					270		

Ser Lys Met Lys Leu Lys Ser Arg Leu Ser Gly Gly Val His Arg Leu
 275 280 285
 Glu Ser Val Glu Glu Tyr Asn Glu Leu Met Val Arg Asn Gly Asp Pro
 290 295 300
 Arg Ile Arg Met Leu Glu Val Ser Arg Asp Gly Arg Lys His Ser Leu
 305 310 315 320
 Pro Gln Leu Leu Asp Ser Thr Gly Thr Ser Gln Glu Tyr His Ile Val
 325 330 335
 Lys Lys Ser Thr Arg Ser Leu Ser Thr Thr His Val Glu Ser Pro Trp
 340 345 350
 Arg Leu Ile Arg Pro Ser Val Ile Ser Ile Ile Gly
 355 360

<210> 65
 <211> 51
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: LRRCT
 Consensus Sequence

<400> 65
 Asn Pro Phe Ile Cys Asp Cys Glu Leu Arg Trp Leu Leu Arg Trp Leu
 1 5 10 15
 Gln Ala Asn Arg His Leu Gln Asp Pro Val Asp Leu Arg Cys Ala Ser
 20 25 30
 Pro Glu Ser Leu Arg Gly Pro Leu Leu Leu Leu Leu Pro Ser Ser Phe
 35 40 45
 Lys Cys Pro
 50

<210> 66
 <211> 51
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: LRRCT
 Consensus Sequence

<400> 66
 Asn Pro Phe Ile Cys Asp Cys Glu Leu Arg Trp Leu Leu Arg Trp Leu
 1 5 10 15
 Arg Glu Pro Arg Arg Leu Glu Asp Pro Glu Asp Leu Arg Cys Ala Ser

20 25 30

Pro Glu Ser Leu Arg Gly Pro Leu Leu Glu Leu Leu Pro Ser Asp Phe
35 40 45

Ser Cys Pro
50

<210> 67
<211> 124
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: CLECT
Consensus Sequence

<400> 67
Cys Pro Ser Gly Trp Val Ser Tyr Pro Gly Gly Lys Cys Tyr Lys Phe
1 5 10 15
Ser Thr Glu Lys Lys Thr Trp Ala Asp Ala Gln Ala Phe Cys Gln Ser
20 25 30
Leu Gly Ala His Leu Ala Ser Ile His Ser Glu Glu Glu Asn Asp Phe
35 40 45
Leu Leu Ser Leu Leu Lys Asn Ser Asn Ser Asp Tyr Tyr Trp Ile Gly
50 55 60
Leu Ser Arg Pro Asp Ser Asn Gly Ser Trp Gln Trp Ser Asp Gly Ser
65 70 75 80
Gly Pro Val Asp Tyr Ser Asn Trp Ala Pro Gly Glu Pro Gly Gly Ser
85 90 95
Gly Asn Cys Val Val Leu Ser Thr Ser Gly Gly Gly Lys Trp Asn Asp
100 105 110
Val Ser Cys Thr Ser Lys Leu Pro Phe Ile Cys Glu
115 120

<210> 68
<211> 114
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: CUB Consensus
Sequence

<400> 68
Cys Gly Gly Thr Leu Thr Ala Ser Ser Gly Thr Ile Thr Ser Pro Asn
1 5 10 15

Tyr Pro Asn Ser Tyr Pro Asn Asn Leu Asn Cys Val Trp Thr Ile Ser
 20 25 30
 Ala Pro Pro Gly Tyr Arg Ile Glu Leu Lys Phe Thr Asp Phe Asp Leu
 35 40 45
 Glu Ser Ser Asp Asn Cys Thr Tyr Asp Tyr Val Glu Ile Tyr Asp Gly
 50 55 60
 Pro Ser Thr Ser Ser Pro Leu Leu Gly Arg Phe Cys Gly Ser Glu Leu
 65 70 75 80
 Pro Pro Pro Ile Ile Ser Ser Ser Ser Asn Ser Met Thr Val Thr Phe
 85 90 95
 Val Ser Asp Ser Ser Val Gln Lys Arg Gly Phe Ser Ala Arg Tyr Ser
 100 105 110

Ala Val

<210> 69
 <211> 110
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: CUB Consensus
 Sequence

<400> 69
 Cys Gly Gly Val Leu Thr Glu Ser Ser Gly Ser Ile Ser Ser Pro Asn
 1 5 10 15
 Tyr Pro Asn Asp Tyr Pro Pro Asn Lys Glu Cys Val Trp Thr Ile Arg
 20 25 30
 Ala Pro Pro Gly Tyr Arg Val Glu Leu Thr Phe Gln Asp Phe Asp Leu
 35 40 45
 Glu Asp His Thr Gly Cys Arg Tyr Asp Tyr Val Glu Ile Arg Asp Gly
 50 55 60
 Asp Gly Ser Ser Ser Pro Leu Leu Gly Lys Phe Cys Gly Ser Gly Pro
 65 70 75 80
 Pro Glu Asp Ile Val Ser Ser Ser Asn Arg Met Thr Ile Lys Phe Val
 85 90 95
 Ser Asp Ala Ser Val Ser Lys Arg Gly Phe Lys Ala Thr Tyr
 100 105 110

<210> 70
 <211> 107
 <212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Lectin C-type
domain Consensus Sequence

<400> 70

Glu	Ser	Lys	Thr	Trp	Ala	Glu	Ala	Gln	Ala	Ala	Cys	Gln	Lys	Leu	Gly
1				5				10						15	
Gly	Gly	Leu	Val	Ser	Ile	Gln	Ser	Ala	Glu	Glu	Gln	Asp	Phe	Leu	Thr
			20					25					30		
Ser	Leu	Thr	Lys	Ala	Ser	Asn	Ser	Tyr	Ala	Trp	Ile	Gly	Leu	Thr	Asp
			35				40					45			
Ile	Asn	Thr	Glu	Gly	Thr	Trp	Val	Trp	Thr	Asp	Gly	Ser	Pro	Val	Asn
	50					55					60				
Tyr	Thr	Asn	Trp	Ala	Pro	Gly	Glu	Pro	Asn	Asn	Arg	Gly	Asn	Lys	Glu
	65				70					75					80
Asp	Cys	Val	Glu	Ile	Tyr	Thr	Asp	Gly	Asn	Lys	Trp	Asn	Asp	Glu	Pro
				85					90					95	
Cys	Gly	Ser	Lys	Leu	Pro	Tyr	Val	Cys	Glu	Phe					
			100					105							

<210> 71

<211> 304

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CNH domain
Consensus Sequence

<400> 71

Tyr	Thr	Ala	Lys	Cys	Asn	His	Pro	Ile	Thr	Cys	Asp	Ala	Lys	Asn	Leu
1				5				10						15	
Leu	Val	Gly	Thr	Glu	Glu	Gly	Leu	Tyr	Val	Leu	Asn	Arg	Ser	Asp	Gln
			20					25					30		
Gly	Gly	Thr	Leu	Glu	Lys	Ile	Ile	Ser	Arg	Arg	Ser	Val	Thr	Gln	Ile
			35				40					45			
Trp	Val	Leu	Glu	Glu	Asn	Asn	Val	Leu	Leu	Met	Ile	Ser	Gly	Lys	Lys
	50					55					60				
Pro	Tyr	Leu	Tyr	Ala	His	Pro	Leu	Ser	Gly	Leu	Arg	Glu	Lys	Asp	Ala
	65				70					75					80
Leu	Gly	Ser	Ala	Arg	Leu	Val	Ile	Arg	Lys	Asn	Val	Trp	Val	Lys	Ile
				85					90					95	

Glu	Asp	Val	Lys	Gly	Cys	His	Leu	Phe	Ala	Val	Val	Asn	Gly	Lys	Arg	100	105	110	
Ile	Leu	Phe	Leu	Cys	Ala	Ala	Leu	Pro	Ser	Ser	Val	Gln	Leu	Leu	Gln	115	120	125	
Trp	Tyr	Asn	Pro	Leu	Lys	Lys	Phe	Lys	Leu	Phe	Lys	Ser	Lys	Phe	Leu	130	135	140	
Lys	Lys	Leu	Ile	Val	Pro	Val	Pro	Leu	Phe	Val	Leu	Leu	Thr	Ser	Ser	145	150	155	160
Ser	Phe	Glu	Leu	Pro	Lys	Ile	Cys	Ile	Gly	Val	Asp	Lys	Asn	Gly	Phe	165	170	175	
Asp	Val	Val	Gln	Phe	His	Gln	Thr	Ser	Leu	Val	Ser	Lys	Glu	Asp	Leu	180	185	190	
Ser	Leu	Pro	Asn	Leu	Asn	Glu	Glu	Thr	Ser	Lys	Lys	Pro	Ile	Pro	Val	195	200	205	
Ile	Gln	Val	Pro	Gln	Ser	Asp	Asp	Glu	Leu	Leu	Leu	Cys	Phe	Asp	Glu	210	215	220	
Phe	Gly	Val	Phe	Val	Asn	Leu	Gln	Gly	Met	Arg	Arg	Ser	Arg	Lys	Pro	225	230	235	240
Ile	Leu	Thr	Trp	Glu	Phe	Met	Pro	Glu	Tyr	Phe	Ala	Tyr	His	Glu	Pro	245	250	255	
Tyr	Leu	Leu	Ala	Phe	His	Ser	Asn	Gly	Ile	Glu	Ile	Arg	Ser	Ile	Glu	260	265	270	
Thr	Gly	Glu	Leu	Leu	Gln	Glu	Leu	Ala	Asp	Arg	Glu	Ala	Arg	Lys	Ile	275	280	285	
Arg	Val	Leu	Gly	Ser	Ser	Asp	Arg	Lys	Ile	Leu	Val	Ser	Ser	Ser	Pro	290	295	300	

<210> 72

<211> 256

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Serine/Threonine protein kinase Consensus
Sequence

<400> 72

Tyr	Glu	Leu	Leu	Glu	Val	Leu	Gly	Lys	Gly	Ala	Phe	Gly	Lys	Val	Tyr	1	5	10	15
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	---	---	----	----

Leu Ala Arg Asp Lys Lys Thr Gly Lys Leu Val Ala Ile Lys Val Ile
 20 25 30
 Lys Lys Glu Lys Leu Lys Lys Lys Lys Arg Glu Arg Ile Leu Arg Glu
 35 40 45
 Ile Lys Ile Leu Lys Lys Leu Asp His Pro Asn Ile Val Lys Leu Tyr
 50 55 60
 Asp Val Phe Glu Asp Asp Asp Lys Leu Tyr Leu Val Met Glu Tyr Cys
 65 70 75 80
 Glu Gly Gly Asp Leu Phe Asp Leu Leu Lys Lys Arg Gly Arg Leu Ser
 85 90 95
 Glu Asp Glu Ala Arg Phe Tyr Ala Arg Gln Ile Leu Ser Ala Leu Glu
 100 105 110
 Tyr Leu His Ser Gln Gly Ile Ile His Arg Asp Leu Lys Pro Glu Asn
 115 120 125
 Ile Leu Leu Asp Ser Asp Gly His Val Lys Leu Ala Asp Phe Gly Leu
 130 135 140
 Ala Lys Gln Leu Asp Ser Gly Gly Thr Leu Leu Thr Thr Phe Val Gly
 145 150 155 160
 Thr Pro Glu Tyr Met Ala Pro Glu Val Leu Leu Gly Lys Gly Tyr Gly
 165 170 175
 Lys Ala Val Asp Ile Trp Ser Leu Gly Val Ile Leu Tyr Glu Leu Leu
 180 185 190
 Thr Gly Lys Pro Pro Phe Pro Gly Asp Asp Gln Leu Leu Ala Leu Phe
 195 200 205
 Lys Lys Ile Gly Lys Pro Pro Pro Pro Phe Pro Pro Pro Glu Trp Lys
 210 215 220
 Ile Ser Pro Glu Ala Lys Asp Leu Ile Lys Lys Leu Leu Val Lys Asp
 225 230 235 240
 Pro Glu Lys Arg Leu Thr Ala Glu Glu Ala Leu Glu His Pro Phe Phe
 245 250 255

<210> 73

<211> 301

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CNH domain
 Consensus Sequence

<400> 73

Asn	Thr	Ala	Lys	Trp	Asn	His	Pro	Ile	Thr	Cys	Asp	Ala	Lys	Ile	Leu	1	5	10	15
Leu	Val	Gly	Thr	Glu	Glu	Gly	Leu	Tyr	Val	Leu	Asn	Ile	Ser	Asp	Gln	20	25	30	
His	Gly	Thr	Leu	Glu	Lys	Leu	Ile	Gly	Arg	Arg	Ser	Val	Thr	Gln	Ile	35	40	45	
Trp	Val	Leu	Glu	Glu	Asn	Asn	Val	Leu	Leu	Met	Ile	Ser	Gly	Lys	Lys	50	55	60	
Pro	Gln	Leu	Tyr	Ser	His	Pro	Leu	Ser	Ala	Leu	Thr	Glu	Lys	Asp	Ala	65	70	75	80
Leu	Gly	Ser	Ala	Arg	Leu	Val	Ile	Arg	Lys	Asn	Val	Leu	Thr	Lys	Ile	85	90	95	
Pro	Asp	Val	Lys	Gly	Cys	His	Leu	Cys	Ala	Val	Val	Asn	Gly	Lys	Arg	100	105	110	
Ile	Leu	Phe	Leu	Cys	His	Ala	Leu	Gln	Ser	Ser	Val	Val	Leu	Leu	Gln	115	120	125	
Trp	Tyr	Asn	Pro	Leu	Lys	Lys	Phe	Lys	Leu	Phe	Lys	Ser	Lys	Phe	Leu	130	135	140	
Phe	Pro	Leu	Ile	Ser	Pro	Val	Pro	Val	Phe	Val	Glu	Leu	Val	Ser	Ser	145	150	155	160
Ser	Phe	Glu	Leu	Pro	Gly	Ile	Cys	Ile	Gly	Ser	Asp	Lys	Asn	Gly	Gly	165	170	175	
Asp	Val	Val	Gln	Phe	His	Gln	Ser	Leu	Val	Ser	Lys	Glu	Asp	Leu	Ser	180	185	190	
Leu	Pro	Phe	Leu	Ser	Glu	Glu	Thr	Ser	Ser	Lys	Pro	Ile	Ser	Val	Val	195	200	205	
Gln	Val	Pro	Ala	Asp	Glu	Leu	Leu	Cys	Tyr	Asp	Glu	Phe	Gly	Val		210	215	220	
Phe	Val	Asn	Leu	Tyr	Gly	Met	Arg	Arg	Ser	Arg	Asn	Pro	Ile	Leu	His	225	230	235	240
Trp	Glu	Phe	Met	Pro	Glu	Ser	Phe	Ala	Tyr	His	Ser	Pro	Tyr	Leu	Leu	245	250	255	
Ala	Phe	His	Asp	Asn	Gly	Ile	Glu	Ile	Arg	Ser	Ile	Lys	Thr	Gly	Glu	260	265	270	
Leu	Leu	Gln	Glu	Leu	Ala	Asp	Arg	Lys	Thr	Arg	Lys	Ile	Arg	Leu	Leu	275	280	285	
Gly	Ser	Ser	Asp	Arg	Lys	Ile	Leu	Leu	Ser	Ser	Ser	Pro							

290

295

300

<210> 74

<211> 256

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Protein kinase
domain Consensus Sequence

<400> 74

Tyr	Glu	Leu	Gly	Glu	Lys	Leu	Gly	Ser	Gly	Ala	Phe	Gly	Lys	Val	Tyr
1				5					10					15	

Lys	Gly	Lys	His	Lys	Asp	Thr	Gly	Glu	Ile	Val	Ala	Ile	Lys	Ile	Leu
			20					25					30		

Lys	Lys	Arg	Ser	Leu	Ser	Glu	Lys	Lys	Lys	Arg	Phe	Leu	Arg	Glu	Ile
		35					40					45			

Gln	Ile	Leu	Arg	Arg	Leu	Ser	His	Pro	Asn	Ile	Val	Arg	Leu	Leu	Gly
	50					55					60				

Val	Phe	Glu	Glu	Asp	Asp	His	Leu	Tyr	Leu	Val	Met	Glu	Tyr	Met	Glu
65					70					75					80

Gly	Gly	Asp	Leu	Phe	Asp	Tyr	Leu	Arg	Arg	Asn	Gly	Leu	Leu	Leu	Ser
				85					90					95	

Glu	Lys	Glu	Ala	Lys	Lys	Ile	Ala	Leu	Gln	Ile	Leu	Arg	Gly	Leu	Glu
			100					105					110		

Tyr	Leu	His	Ser	Arg	Gly	Ile	Val	His	Arg	Asp	Leu	Lys	Pro	Glu	Asn
		115					120					125			

Ile	Leu	Leu	Asp	Glu	Asn	Gly	Thr	Val	Lys	Ile	Ala	Asp	Phe	Gly	Leu
	130					135					140				

Ala	Arg	Lys	Leu	Glu	Ser	Ser	Ser	Tyr	Glu	Lys	Leu	Thr	Thr	Phe	Val
145					150					155					160

Gly	Thr	Pro	Glu	Tyr	Met	Ala	Pro	Glu	Val	Leu	Glu	Gly	Arg	Gly	Tyr
			165						170					175	

Ser	Ser	Lys	Val	Asp	Val	Trp	Ser	Leu	Gly	Val	Ile	Leu	Tyr	Glu	Leu
			180					185					190		

Leu	Thr	Gly	Lys	Leu	Pro	Phe	Pro	Gly	Ile	Asp	Pro	Leu	Glu	Glu	Leu
		195						200				205			

Phe	Arg	Ile	Lys	Glu	Arg	Pro	Arg	Leu	Arg	Leu	Pro	Leu	Pro	Pro	Asn
	210					215					220				

Cys	Ser	Glu	Glu	Leu	Lys	Asp	Leu	Ile	Lys	Lys	Cys	Leu	Asn	Lys	Asp
225					230					235					240

Pro Glu Lys Arg Pro Thr Ala Lys Glu Ile Leu Asn His Pro Trp Phe
245 250 255

<210> 75

<211> 258

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Tyrosine
Kinase Consensus Sequence

<400> 75

Leu Thr Leu Gly Lys Lys Leu Gly Glu Gly Ala Phe Gly Glu Val Tyr
1 5 10 15

Lys Gly Thr Leu Lys Gly Lys Gly Gly Val Glu Val Glu Val Ala Val
20 25 30

Lys Thr Leu Lys Glu Asp Ala Ser Glu Gln Gln Ile Glu Glu Phe Leu
35 40 45

Arg Glu Ala Arg Leu Met Arg Lys Leu Asp His Pro Asn Ile Val Lys
50 55 60

Leu Leu Gly Val Cys Thr Glu Glu Glu Pro Leu Met Ile Val Met Glu
65 70 75 80

Tyr Met Glu Gly Gly Asp Leu Leu Asp Tyr Leu Arg Lys Asn Arg Pro
85 90 95

Lys Glu Leu Ser Leu Ser Asp Leu Leu Ser Phe Ala Leu Gln Ile Ala
100 105 110

Arg Gly Met Glu Tyr Leu Glu Ser Lys Asn Phe Val His Arg Asp Leu
115 120 125

Ala Ala Arg Asn Cys Leu Val Gly Glu Asn Lys Thr Val Lys Ile Ala
130 135 140

Asp Phe Gly Leu Ala Arg Asp Leu Tyr Asp Asp Asp Tyr Tyr Arg Lys
145 150 155 160

Lys Lys Ser Pro Arg Leu Pro Ile Arg Trp Met Ala Pro Glu Ser Leu
165 170 175

Lys Asp Gly Lys Phe Thr Ser Lys Ser Asp Val Trp Ser Phe Gly Val
180 185 190

Leu Leu Trp Glu Ile Phe Thr Leu Gly Glu Ser Pro Tyr Pro Gly Met
195 200 205

Ser Asn Glu Glu Val Leu Glu Tyr Leu Lys Lys Gly Tyr Arg Leu Pro
 210 215 220

Gln Pro Pro Asn Cys Pro Asp Glu Ile Tyr Asp Leu Met Leu Gln Cys
 225 230 235 240

Trp Ala Glu Asp Pro Glu Asp Arg Pro Thr Phe Ser Glu Leu Val Glu
 245 250 255

Arg Leu

<210> 76

<211> 430

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Sema domain
 Consensus Sequence

<400> 76

Leu Gln Asn Leu Leu Leu Asp Glu Asp Asn Gly Thr Leu Tyr Val Gly
 1 5 10 15

Ala Arg Asn Arg Leu Tyr Val Leu Ser Leu Asn Leu Ile Ser Glu Ala
 20 25 30

Glu Val Lys Thr Gly Pro Val Leu Ser Ser Pro Asp Cys Glu Glu Cys
 35 40 45

Val Ser Lys Gly Lys Asp Pro Pro Thr Asp Cys Val Asn Phe Ile Arg
 50 55 60

Leu Leu Leu Asp Tyr Asn Ala Asp His Leu Leu Val Cys Gly Thr Asn
 65 70 75 80

Ala Phe Gln Pro Val Cys Arg Leu Ile Asn Leu Gly Asn Leu Asp Arg
 85 90 95

Leu Glu Val Gly Arg Glu Ser Gly Arg Gly Arg Cys Pro Phe Asp Pro
 100 105 110

Gln His Asn Ser Thr Ala Val Leu Val Asp Gly Glu Leu Tyr Val Gly
 115 120 125

Thr Val Ala Asp Phe Ser Gly Ser Asp Pro Ala Ile Tyr Arg Ser Leu
 130 135 140

Ser Val Arg Arg Leu Lys Gly Thr Ser Gly Pro Ser Leu Arg Thr Val
 145 150 155 160

Leu Tyr Asp Ser Arg Trp Leu Asn Glu Pro Asn Phe Val Tyr Ala Phe
 165 170 175

Glu Ser Gly Asp Phe Val Tyr Phe Phe Phe Arg Glu Thr Ala Val Glu

180					185					190						
Asp	Glu	Asn	Cys	Gly	Lys	Ala	Val	Val	Ser	Arg	Val	Ala	Arg	Val	Cys	
195					200					205						
Lys	Asn	Asp	Val	Gly	Gly	Pro	Arg	Ser	Leu	Ser	Lys	Lys	Trp	Thr	Ser	
210					215					220						
Phe	Leu	Lys	Ala	Arg	Leu	Glu	Cys	Ser	Val	Pro	Gly	Glu	Phe	Pro	Phe	
225					230					235					240	
Tyr	Phe	Asn	Glu	Leu	Gln	Ala	Ala	Phe	Leu	Leu	Pro	Ala	Gly	Ser	Glu	
245					250					255						
Ser	Asp	Asp	Val	Leu	Tyr	Gly	Val	Phe	Ser	Thr	Ser	Ser	Asn	Pro	Ile	
260					265					270						
Pro	Gly	Ser	Ala	Val	Cys	Ala	Phe	Ser	Leu	Ser	Asp	Ile	Asn	Ala	Val	
275					280					285						
Phe	Asn	Glu	Pro	Phe	Lys	Glu	Cys	Glu	Thr	Gly	Asn	Ser	Gln	Trp	Leu	
290					295					300						
Pro	Tyr	Pro	Arg	Gly	Leu	Val	Pro	Phe	Pro	Arg	Pro	Gly	Thr	Cys	Pro	
305					310					315					320	
Asn	Thr	Pro	Leu	Ser	Ser	Lys	Asp	Leu	Pro	Asp	Asp	Val	Leu	Asn	Phe	
325					330					335						
Ile	Lys	Thr	His	Pro	Leu	Met	Asp	Glu	Val	Val	Gln	Pro	Leu	Thr	Gly	
340					345					350						
Arg	Pro	Leu	Phe	Val	Lys	Thr	Asp	Ser	Asn	Tyr	Leu	Leu	Thr	Ser	Ile	
355					360					365						
Ala	Val	Asp	Arg	Val	Arg	Thr	Asp	Gly	Gly	Asn	Tyr	Thr	Val	Leu	Phe	
370					375					380						
Leu	Gly	Thr	Ser	Asp	Gly	Arg	Ile	Leu	Lys	Val	Val	Leu	Ser	Arg	Ser	
385					390					395					400	
Ser	Ser	Ser	Ser	Glu	Ser	Val	Val	Leu	Glu	Glu	Ile	Ser	Val	Phe	Asp	
405					410					415						
Pro	Gly	Ser	Pro	Val	Ser	Asp	Leu	Val	Leu	Ser	Pro	Lys	Lys			
420					425					430						

<210> 77

<211> 433

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Sema domain
Consensus Sequence

<400> 77

Phe	Val	Thr	Leu	Leu	Leu	Asp	Glu	Asp	Arg	Gly	Arg	Leu	Tyr	Val	Gly	1	5	10	15
Ala	Arg	Asn	Arg	Val	Tyr	Val	Leu	Asn	Leu	Glu	Asp	Leu	Ser	Glu	Val	20	25	30	
Leu	Asn	Leu	Lys	Thr	Gly	Trp	Pro	Gly	Ser	Cys	Glu	Thr	Cys	Glu	Glu	35	40	45	
Cys	Asn	Met	Lys	Gly	Lys	Ser	Pro	Leu	Thr	Glu	Cys	Thr	Asn	Phe	Ile	50	55	60	
Arg	Val	Leu	Gln	Ala	Tyr	Asn	Asp	Thr	His	Leu	Tyr	Val	Cys	Gly	Thr	65	70	75	80
Asn	Ala	Phe	Gln	Pro	Val	Cys	Thr	Leu	Ile	Asn	Leu	Gly	Asp	Leu	Phe	85	90	95	
Ser	Leu	Asp	Val	Asp	Asn	Glu	Glu	Asp	Gly	Cys	Gly	Asp	Cys	Pro	Tyr	100	105	110	
Asp	Pro	Leu	Gly	Asn	Thr	Thr	Ser	Val	Leu	Val	Gln	Gly	Gly	Glu	Leu	115	120	125	
Tyr	Ser	Gly	Thr	Val	Ile	Asp	Phe	Ser	Gly	Arg	Asp	Pro	Ser	Ile	Arg	130	135	140	
Arg	Leu	Leu	Gly	Ser	His	Asp	Gly	Leu	Arg	Thr	Glu	Phe	His	Asp	Ser	145	150	155	160
Lys	Trp	Leu	Asn	Leu	Pro	Asn	Phe	Val	Asp	Ser	Tyr	Pro	Ile	His	Tyr	165	170	175	
Val	His	Ser	Phe	Ser	Asp	Asp	Lys	Val	Tyr	Phe	Phe	Phe	Arg	Glu	Thr	180	185	190	
Ala	Val	Glu	Asp	Ser	Asn	Cys	Lys	Thr	Ile	His	Ser	Arg	Val	Ala	Arg	195	200	205	
Val	Cys	Lys	Asn	Asp	Pro	Gly	Gly	Arg	Ser	Tyr	Leu	Glu	Leu	Asn	Lys	210	215	220	
Trp	Thr	Thr	Phe	Leu	Lys	Ala	Arg	Leu	Asn	Cys	Ser	Ile	Pro	Gly	Glu	225	230	235	240
Gly	Thr	Pro	Phe	Tyr	Phe	Asn	Glu	Leu	Gln	Ala	Ala	Phe	Val	Leu	Pro	245	250	255	
Thr	Gly	Ala	Asp	Thr	Asp	Pro	Val	Leu	Tyr	Gly	Val	Phe	Thr	Thr	Ser	260	265	270	
Ser	Asn	Ser	Ser	Ala	Gly	Ser	Ala	Val	Cys	Ala	Phe	Ser	Met	Ser	Asp	275	280	285	
Ile	Asn	Gln	Val	Phe	Glu	Gly	Pro	Phe	Lys	His	Gln	Ser	Pro	Asn	Ser	290	295	300	

Lys Trp Leu Pro Tyr Arg Gly Lys Val Pro Gln Pro Arg Pro Gly Gln
 305 310 315 320
 Cys Pro Asn Ala Ser Gly Leu Asn Leu Pro Asp Asp Thr Leu Asn Phe
 325 330 335
 Ile Arg Cys His Pro Leu Met Asp Glu Val Val Pro Pro Leu His Asn
 340 345 350
 Val Pro Leu Phe Val Gly Gln Ser Gly Asn Tyr Arg Leu Thr Ser Ile
 355 360 365
 Ala Val Asp Arg Val Arg Ala Gly Asp Gly Gln Ile Tyr Thr Val Leu
 370 375 380
 Phe Leu Gly Thr Asp Asp Gly Arg Val Leu Lys Gln Val Val Leu Ser
 385 390 395 400
 Arg Ser Ser Ser Ala Ser Tyr Leu Val Val Val Leu Glu Glu Ser Leu
 405 410 415
 Val Phe Pro Asp Gly Glu Pro Val Gln Arg Met Val Ile Ser Ser Lys
 420 425 430

Asn

<210> 78

<211> 85

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: TIG domain
Consensus Sequence

<400> 78

Pro Val Ile Thr Ser Ile Ser Pro Ser Ser Gly Pro Leu Ser Gly Gly
 1 5 10 15
 Thr Glu Ile Thr Ile Thr Gly Ser Asn Leu Gly Ser Gly Glu Asp Ile
 20 25 30
 Lys Val Thr Phe Gly Gly Thr Glu Cys Asp Val Val Ser Gln Glu Ala
 35 40 45
 Ser Gln Ile Val Cys Lys Thr Pro Pro Tyr Ala Asn Gly Gly Pro Gln
 50 55 60
 Pro Val Thr Val Ser Leu Asp Gly Gly Gly Leu Ser Ser Ser Pro Val
 65 70 75 80
 Thr Phe Thr Tyr Val
 85

<210> 79
 <211> 85
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: TIG Consensus
 Sequence

<400> 79
 Pro Val Ile Thr Ser Ile Ser Pro Ser Ser Gly Pro Leu Ser Gly Gly
 1 5 10 15
 Thr Glu Ile Thr Ile Thr Gly Ser Asn Leu Gly Ser Gly Glu Asp Ile
 20 25 30
 Lys Val Thr Phe Gly Gly Thr Glu Cys Asp Val Val Ser Gln Glu Ala
 35 40 45
 Ser Gln Ile Val Cys Lys Thr Pro Pro Tyr Ala Asn Gly Gly Pro Gln
 50 55 60
 Pro Val Thr Val Ser Leu Asp Gly Gly Gly Leu Ser Ser Ser Pro Val
 65 70 75 80
 Thr Phe Thr Tyr Val
 85

<210> 80
 <211> 85
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: TIG Consensus
 Sequence

<400> 80
 Pro Val Ile Thr Ser Ile Ser Pro Ser Ser Gly Pro Leu Ser Gly Gly
 1 5 10 15
 Thr Glu Ile Thr Ile Thr Gly Ser Asn Leu Gly Ser Gly Glu Asp Ile
 20 25 30
 Lys Val Thr Phe Gly Gly Thr Glu Cys Asp Val Val Ser Gln Glu Ala
 35 40 45
 Ser Gln Ile Val Cys Lys Thr Pro Pro Tyr Ala Asn Gly Gly Pro Gln
 50 55 60
 Pro Val Thr Val Ser Leu Asp Gly Gly Gly Leu Ser Ser Ser Pro Val
 65 70 75 80
 Thr Phe Thr Tyr Val
 85

<210> 81
 <211> 47
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PSI domain
 Consensus Sequence

<400> 81
 Arg Cys Ser Ala Tyr Thr Ser Cys Ser Glu Cys Leu Leu Ala Arg Asp
 1 5 10 15
 Pro Tyr Cys Ala Trp Cys Ser Ser Gln Gly Arg Cys Thr Ser Gly Glu
 20 25 30
 Arg Cys Asp Ser Leu Arg Gln Asn Trp Ser Ser Gly Gln Cys Pro
 35 40 45

<210> 82
 <211> 93
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: IPT Consensus
 Sequence

<400> 82
 Asp Pro Val Ile Thr Arg Ile Ser Pro Asn Ser Gly Pro Leu Ser Gly
 1 5 10 15
 Gly Thr Arg Ile Thr Leu Cys Gly Lys Asn Leu Asp Ser Ile Ser Val
 20 25 30
 Val Phe Val Glu Val Gly Val Gly Glu Val Pro Cys Thr Phe Leu Pro
 35 40 45
 Ser Asp Val Ser Gln Thr Ala Ile Val Cys Lys Thr Pro Pro Tyr His
 50 55 60
 Asn Ile Pro Gly Ser Val Pro Val Arg Val Glu Val Gly Leu Arg Asn
 65 70 75 80
 Gly Gly Val Pro Gly Glu Pro Ser Pro Phe Thr Tyr Val
 85 90

<210> 83
 <211> 254
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: 7
transmembrane receptor Consensus Sequence

<400> 83

Gly	Asn	Leu	Leu	Val	Ile	Leu	Val	Ile	Leu	Arg	Thr	Lys	Lys	Leu	Arg	1	5	10	15
Thr	Pro	Thr	Asn	Ile	Phe	Leu	Leu	Asn	Leu	Ala	Val	Ala	Asp	Leu	Leu	20	25	30	
Phe	Leu	Leu	Thr	Leu	Pro	Pro	Trp	Ala	Leu	Tyr	Tyr	Leu	Val	Gly	Gly	35	40	45	
Asp	Trp	Val	Phe	Gly	Asp	Ala	Leu	Cys	Lys	Leu	Val	Gly	Ala	Leu	Phe	50	55	60	
Val	Val	Asn	Gly	Tyr	Ala	Ser	Ile	Leu	Leu	Leu	Thr	Ala	Ile	Ser	Ile	65	70	75	80
Asp	Arg	Tyr	Leu	Ala	Ile	Val	His	Pro	Leu	Arg	Tyr	Arg	Arg	Ile	Arg	85	90	95	
Thr	Pro	Arg	Arg	Ala	Lys	Val	Leu	Ile	Leu	Leu	Val	Trp	Val	Leu	Ala	100	105	110	
Leu	Leu	Leu	Ser	Leu	Pro	Pro	Leu	Leu	Phe	Ser	Trp	Leu	Arg	Thr	Val	115	120	125	
Glu	Glu	Gly	Asn	Thr	Thr	Val	Cys	Leu	Ile	Asp	Phe	Pro	Glu	Glu	Ser	130	135	140	
Val	Lys	Arg	Ser	Tyr	Val	Leu	Leu	Ser	Thr	Leu	Val	Gly	Phe	Val	Leu	145	150	155	160
Pro	Leu	Leu	Val	Ile	Leu	Val	Cys	Tyr	Thr	Arg	Ile	Leu	Arg	Thr	Leu	165	170	175	
Arg	Lys	Arg	Ala	Arg	Ser	Gln	Arg	Ser	Leu	Lys	Arg	Arg	Ser	Ser	Ser	180	185	190	
Glu	Arg	Lys	Ala	Ala	Lys	Met	Leu	Leu	Val	Val	Val	Val	Val	Phe	Val	195	200	205	
Leu	Cys	Trp	Leu	Pro	Tyr	His	Ile	Val	Leu	Leu	Leu	Asp	Ser	Leu	Cys	210	215	220	
Leu	Leu	Ser	Ile	Trp	Arg	Val	Leu	Pro	Thr	Ala	Leu	Leu	Ile	Thr	Leu	225	230	235	240
Trp	Leu	Ala	Tyr	Val	Asn	Ser	Cys	Leu	Asn	Pro	Ile	Ile	Tyr	245	250				

<210> 84

<211> 402

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: ANF receptor
Consensus Sequence

<400> 84

Gly	Leu	Ile	Asn	Tyr	Ala	Val	Arg	Gly	Ile	Thr	Arg	Leu	Glu	Ala	Met
1				5					10					15	
Leu	Gly	Ala	Phe	Asp	Arg	Ile	Asn	Ala	Asp	Pro	Ala	Leu	Leu	Pro	Gly
			20					25					30		
Leu	Ala	Leu	Gly	Leu	Ala	Ile	Leu	Asp	Ile	Asn	Ser	Leu	Arg	Asn	Val
		35					40					45			
Ala	Leu	Glu	Gln	Ser	Phe	Thr	Phe	Val	Tyr	Gly	Leu	Leu	Ile	Lys	Cys
	50					55					60				
Asp	Cys	Ser	Ser	Val	Arg	Cys	Ala	Gly	Gly	Asp	Leu	Ala	Leu	Thr	His
65					70					75					80
Gly	Val	Ala	Gly	Val	Ile	Gly	Pro	Ser	Cys	Ser	Ser	Ser	Ala	Ile	Gln
				85					90					95	
Val	Ala	Asn	Leu	Ala	Ser	Leu	Leu	Asn	Ile	Pro	Met	Ile	Ser	Tyr	Gly
		100						105					110		
Ser	Thr	Ala	Pro	Glu	Leu	Ser	Asp	Lys	Thr	Arg	Tyr	Pro	Thr	Phe	Ser
		115					120					125			
Arg	Thr	Ile	Pro	Ser	Asp	Ala	Phe	Gln	Gly	Leu	Ala	Met	Val	Asp	Ile
	130					135					140				
Phe	Lys	His	Phe	Asn	Trp	Asn	Tyr	Val	Ser	Val	Val	Tyr	Ser	Asp	Gly
145					150					155					160
Thr	Tyr	Gly	Glu	Glu	Gly	Cys	Glu	Ala	Phe	Ile	Glu	Ala	Leu	Arg	Ala
				165					170				175		
Arg	Gly	Gly	Cys	Ile	Ala	Leu	Ser	Val	Lys	Ile	Gly	Glu	Phe	Asp	Arg
			180					185					190		
Gly	Asp	Glu	Glu	Asp	Phe	Asp	Lys	Leu	Leu	Arg	Glu	Leu	Lys	Arg	Arg
	195						200					205			
Ala	Arg	Val	Val	Val	Met	Cys	Gly	His	Gly	Glu	Thr	Leu	Arg	Glu	Leu
	210					215					220				
Leu	Glu	Ala	Ala	Leu	Arg	Leu	Gly	Leu	Thr	Gly	Glu	Asp	Tyr	Val	Phe
225					230					235					240
Ile	Ser	Asp	Asp	Leu	Phe	Asn	Lys	Ser	Leu	Pro	Ala	Glu	Pro	Gly	Ala
				245					250					255	
Pro	Gly	Ala	Ile	Glu	Leu	Ala	Asn	Ala	Ser	Met	Leu	Arg	Phe	Ala	Tyr
		260					265						270		

Tyr Phe Val Leu Val Leu Thr Leu Asn Asn Pro Arg Asn Pro Trp Phe
 275 280 285
 Leu Glu Phe Trp Lys Glu Asn Phe Ile Cys Ala Leu Gln Asp Phe Leu
 290 295 300
 Thr Leu Glu Pro Tyr Glu Gln Glu Gly Lys Ala Gly Phe Val Tyr Asp
 305 310 315 320
 Ala Val Tyr Leu Tyr Ala His Ala Leu His Asn Thr Thr Leu Ala Leu
 325 330 335
 Gly Gly Ser Trp Val Asp Gly Glu Lys Leu Val Gln His Leu Arg Asn
 340 345 350
 Leu Thr Phe Glu Gly Val Thr Gly Pro Val Thr Phe Asp Glu Asn Gly
 355 360 365
 Asp Arg Asp Gly Asp Tyr Val Leu Leu Asp Thr Gln Asn Thr Glu Thr
 370 375 380
 Gly Gln Leu Lys Val Thr Gly Thr Tyr Asp Gly Val Gly Lys Trp Thr
 385 390 395 400
 Glu Pro

<210> 85

<211> 256

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: 7
 transmembrane receptor (metabotropic E family)
 Consensus Sequence

<400> 85

Leu Gly Ile Val Leu Val Ala Leu Ala Val Leu Gly Ile Val Leu Thr
 1 5 10 15
 Leu Phe Val Leu Val Val Phe Val Lys His Arg Asp Thr Pro Ile Val
 20 25 30
 Lys Ala Ser Asn Arg Glu Leu Ser Tyr Leu Leu Leu Ile Gly Leu Ile
 35 40 45
 Leu Cys Tyr Leu Cys Ser Phe Leu Phe Ile Gly Lys Pro Ser Glu Thr
 50 55 60
 Ser Cys Ile Leu Arg Arg Ile Leu Phe Gly Leu Gly Phe Thr Leu Cys
 65 70 75 80
 Tyr Ser Ala Leu Leu Ala Lys Thr Asn Arg Val Leu Arg Ile Phe Arg
 85 90 95

Ala Lys Lys Pro Gly Ser Gly Lys Pro Lys Phe Ile Ser Pro Trp Ala
 100 105 110
 Gln Val Leu Ile Val Leu Ile Leu Val Leu Ile Gln Val Ile Ile Cys
 115 120 125
 Val Ile Trp Leu Val Val Glu Pro Pro Arg Pro Thr Ile Asp Ile Tyr
 130 135 140
 Ser Glu Lys Glu Lys Ile Ile Leu Glu Cys Asn Lys Gly Ser Met Val
 145 150 155 160
 Ala Phe Val Val Val Leu Gly Tyr Asp Gly Leu Leu Ala Val Leu Cys
 165 170 175
 Thr Phe Leu Ala Phe Leu Thr Arg Asn Leu Pro Glu Asn Phe Asn Glu
 180 185 190
 Ala Lys Phe Ile Gly Phe Ser Met Leu Thr Phe Cys Ile Val Trp Val
 195 200 205
 Ala Phe Ile Pro Ile Tyr Leu Ser Thr Pro Gly Lys Val Gln Val Ala
 210 215 220
 Val Glu Ile Phe Ser Ile Leu Ala Ser Ser Thr Val Leu Leu Gly Cys
 225 230 235 240
 Leu Phe Val Pro Lys Cys Tyr Ile Ile Leu Phe Arg Pro Glu Lys Asn
 245 250 255

<210> 86
 <211> 86
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PDZ domain
 Consensus Sequence

<400> 86
 Glu Pro Arg Leu Val Glu Leu Glu Lys Gly Gly Gly Gly Leu Gly Phe
 1 5 10 15
 Ser Leu Val Gly Gly Lys Asp Ser Gly Asp Gly Gly Val Val Val Ser
 20 25 30
 Ser Val Val Pro Gly Ser Pro Ala Ala Lys Ala Gly Leu Lys Pro Gly
 35 40 45
 Asp Val Ile Leu Glu Val Asn Gly Thr Ser Val Glu Gly Leu Thr His
 50 55 60
 Leu Glu Ala Val Asp Leu Leu Lys Glu Ala Gly Gly Lys Val Thr Leu

65	70	75	80
Thr Val Leu Arg Gly Gly			
85			
<210> 87			
<211> 25			
<212> DNA			
<213> Artificial Sequence			
<220>			
<223> Description of Artificial Sequence: NOV1c Primer 1			
<400> 87			
tcatcacatg acaacatgaa gctgt			25
<210> 88			
<211> 26			
<212> DNA			
<213> Artificial Sequence			
<220>			
<223> Description of Artificial Sequence: NOV1c Primer 2			
<400> 88			
gaaagccctc aaactctcca tctatg			26
<210> 89			
<211> 22			
<212> DNA			
<213> Artificial Sequence			
<220>			
<223> Description of Artificial Sequence: NOV7a Primer 1			
<400> 89			
ccaatctctg atgccctgcg at			22
<210> 90			
<211> 20			
<212> DNA			
<213> Artificial Sequence			
<220>			
<223> Description of Artificial Sequence: NOV7a Primer 2			
<400> 90			
aggtcagtgc cggagcctcc			20
<210> 91			
<211> 19			
<212> DNA			

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag273b Forward

<400> 91
cggttgacg atgcttcac 19

<210> 92
<211> 32
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag273b Probe

<400> 92
tgacttttct gggcttacca atgctatttc aa 32

<210> 93
<211> 27
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag273b Reverse

<400> 93
gcacctatct caatatctgc aatattg 27

<210> 94
<211> 20
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag1094 Forward

<400> 94
atggactgga aaacctggaa 20

<210> 95
<211> 29
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag1094 Probe

<400> 95
tcctgcaagc agataacaat tttatcaca 29

<210> 96
 <211> 20
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Ag1094 Reverse

 <400> 96
 tgctaaaggc acttggttca 20

 <210> 97
 <211> 21
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Ag2100 Forward

 <400> 97
 agatccctgg aacagaggat t 21

 <210> 98
 <211> 26
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Ag2100 Probe

 <400> 98
 tgtctgaagc caataaactt gcagca 26

 <210> 99
 <211> 21
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Ag2100 Reverse

 <400> 99
 ccttcatgtt cctttgggta a 21

 <210> 100
 <211> 22
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Ag217 Forward

 <400> 100

atctgtgctg aggcattgtc ct 22

<210> 101
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag217 Probe

<400> 101
atcctcctcc ctccccggct ctc 23

<210> 102
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag217 Reverse

<400> 102
ctgcatggct ggtgtgatg 19

<210> 103
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag850 Forward

<400> 103
cctttcttct cttcctcctc aa 22

<210> 104
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag850 Probe

<400> 104
cacctggcga gtgctcctct ctg 23

<210> 105
<211> 19
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag850 Reverse

<400> 105
 ggtggatggc gttgtagag 19

<210> 106
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Ag1469 Forward

<400> 106
 cgtacgtctt ccatgatgag tt 22

<210> 107
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Ag1469 Probe

<400> 107
 cgtggcctcg atgattaaga tccctt 26

<210> 108
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Ag1469 Reverse

<400> 108
 aagtcaggga tgatggtgaa g 21

<210> 109
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Ag2976 Forward

<400> 109
 accccaaatg gattccatta 20

<210> 110
 <211> 26
 <212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag2976 Probe

<400> 110
ccctcatgga tctgcataac cacaca 26

<210> 111
<211> 20
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag2976 Reverse

<400> 111
cttgtgtgtg catgcttgtc 20

<210> 112
<211> 22
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag760 Forward

<400> 112
caccatgaca acgacaccta ta 22

<210> 113
<211> 25
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag760 Probe

<400> 113
atatggcacc aacatcacat gcacg 25

<210> 114
<211> 22
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag760 Reverse

<400> 114
tgggtagaaa gtgtgtgtga aa 22

<210> 115
 <211> 22
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Ag1537 Forward

 <400> 115
 aaggagctgg aagagaagaa ga 22

 <210> 116
 <211> 26
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Ag1537 Probe

 <400> 116
 atcagaaact cagccctgga cacctg 26

 <210> 117
 <211> 19
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Ag1537 Reverse

 <400> 117
 gctgcgactt ggtcttgat 19

 <210> 118
 <211> 22
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Ag147 Forward

 <400> 118
 tgaagacagc acctccctat ca 22

 <210> 119
 <211> 21
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Ag147 Probe

 <400> 119

cggtccgtg ctgtcaccca g 21

<210> 120
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag147 Reverse

<400> 120
aagaatcctc agcatcgcca ta 22

<210> 121
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag718 Forward

<400> 121
agaaggaatc tctgggaaag ct 22

<210> 122
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag718 Probe

<400> 122
ccactggaga tgcttggtgc tctacca 27

<210> 123
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag718 Reverse

<400> 123
gacagagcac tggctagttc ac 22

<210> 124
<211> 22
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag3681/Ag4085
Forward

<400> 124
gaatcatcca caagtcattcc at 22

<210> 125
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag3681/Ag4085
Probe

<400> 125
ctcactccca tctcatgcct cccag 25

<210> 126
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag3681/Ag4085
Reverse

<400> 126
catgctgaat gatcgtgaca 20

<210> 127
<211> 1011
<212> PRT
<213> Homo sapiens

<400> 127
Ile Ser Asn Ser Ser Asp Thr Val Glu Cys Glu Cys Ser Glu Asn Trp
1 5 10 15
Lys Gly Glu Ala Cys Asp Ile Pro His Cys Thr Asp Asn Cys Gly Phe
20 25 30
Pro His Arg Gly Ile Cys Asn Ser Ser Asp Val Arg Gly Cys Ser Cys
35 40 45
Phe Ser Asp Trp Gln Gly Pro Gly Cys Ser Val Pro Val Pro Ala Asn
50 55 60
Gln Ser Phe Trp Thr Arg Glu Glu Tyr Ser Asn Leu Lys Leu Pro Arg
65 70 75 80
Ala Ser His Lys Ala Val Val Asn Gly Asn Ile Met Trp Val Val Gly
85 90 95

Pro Gly Ile Arg Cys Val Trp Asn Thr Gly Ser Ser Gln Cys Ile Ser
 405 410 415
 Trp Ala Leu Ala Thr Asp Glu Gln Glu Glu Lys Leu Lys Ser Glu Cys
 420 425 430
 Phe Ser Lys Arg Thr Leu Asp His Asp Arg Cys Asp Gln His Thr Asp
 435 440 445
 Cys Tyr Ser Cys Thr Ala Asn Thr Asn Asp Cys His Trp Cys Asn Asp
 450 455 460
 His Cys Val Pro Arg Asn His Ser Cys Ser Glu Gly Gln Ile Ser Ile
 465 470 475 480
 Phe Arg Tyr Glu Asn Cys Pro Lys Asp Asn Pro Met Tyr Tyr Cys Asn
 485 490 495
 Lys Lys Thr Ser Cys Arg Ser Cys Ala Leu Asp Gln Asn Cys Gln Trp
 500 505 510
 Glu Pro Arg Asn Gln Glu Cys Ile Ala Leu Pro Glu Asn Ile Cys Gly
 515 520 525
 Ile Gly Trp His Leu Val Gly Asn Ser Cys Leu Lys Ile Thr Thr Ala
 530 535 540
 Lys Glu Asn Tyr Asp Asn Ala Lys Leu Phe Cys Arg Asn His Asn Ala
 545 550 555 560
 Leu Leu Ala Ser Leu Thr Thr Gln Lys Lys Val Glu Phe Val Leu Lys
 565 570 575
 Gln Leu Arg Ile Met Gln Ser Ser Gln Ser Met Ser Lys Leu Thr Leu
 580 585 590
 Thr Pro Trp Val Gly Leu Arg Lys Ile Asn Val Ser Tyr Trp Cys Trp
 595 600 605
 Glu Asp Met Ser Pro Phe Thr Asn Ser Leu Leu Gln Trp Met Pro Ser
 610 615 620
 Glu Pro Ser Asp Ala Gly Phe Cys Gly Ile Leu Ser Glu Pro Ser Thr
 625 630 635 640
 Arg Gly Leu Lys Ala Ala Thr Cys Ile Asn Pro Leu Asn Gly Ser Val
 645 650 655
 Cys Glu Arg Pro Ala Asn His Ser Ala Lys Gln Cys Arg Thr Pro Cys
 660 665 670
 Ala Leu Arg Thr Ala Cys Gly Asp Cys Thr Ser Gly Ser Ser Glu Cys
 675 680 685
 Met Trp Cys Ser Asn Met Lys Gln Cys Val Asp Ser Asn Ala Tyr Val
 690 695 700

Ala	Ser	Phe	Pro	Phe	Gly	Gln	Cys	Met	Glu	Trp	Tyr	Thr	Met	Ser	Thr	
705					710					715					720	
Cys	Pro	Pro	Glu	Asn	Cys	Ser	Gly	Tyr	Cys	Thr	Cys	Ser	His	Cys	Leu	
				725					730					735		
Glu	Gln	Pro	Gly	Cys	Gly	Trp	Cys	Thr	Asp	Pro	Ser	Asn	Thr	Gly	Lys	
			740					745					750			
Gly	Lys	Cys	Ile	Glu	Gly	Ser	Tyr	Lys	Gly	Pro	Val	Lys	Met	Pro	Ser	
		755					760					765				
Gln	Ala	Pro	Thr	Gly	Asn	Phe	Tyr	Pro	Gln	Pro	Leu	Leu	Asn	Ser	Ser	
	770					775					780					
Met	Cys	Leu	Glu	Asp	Ser	Arg	Tyr	Asn	Trp	Ser	Phe	Ile	His	Cys	Pro	
785					790					795					800	
Ala	Cys	Gln	Cys	Asn	Gly	His	Ser	Lys	Cys	Ile	Asn	Gln	Ser	Ile	Cys	
				805					810						815	
Glu	Lys	Cys	Glu	Asn	Leu	Thr	Thr	Gly	Lys	His	Cys	Glu	Thr	Cys	Ile	
			820					825					830			
Ser	Gly	Phe	Tyr	Gly	Asp	Pro	Thr	Asn	Gly	Gly	Lys	Cys	Gln	Pro	Cys	
		835					840					845				
Lys	Cys	Asn	Gly	His	Ala	Ser	Leu	Cys	Asn	Thr	Asn	Thr	Gly	Lys	Cys	
	850					855				860						
Phe	Cys	Thr	Thr	Lys	Gly	Val	Lys	Gly	Asp	Glu	Cys	Gln	Leu	Cys	Glu	
865					870					875					880	
Val	Glu	Asn	Arg	Tyr	Gln	Gly	Asn	Pro	Leu	Arg	Gly	Thr	Cys	Tyr	Tyr	
				885					890					895		
Thr	Leu	Leu	Ile	Asp	Tyr	Gln	Phe	Thr	Phe	Ser	Leu	Ser	Gln	Glu	Asp	
			900					905					910			
Asp	Arg	Tyr	Tyr	Thr	Ala	Ile	Asn	Phe	Val	Ala	Thr	Pro	Asp	Glu	Gln	
		915					920					925				
Asn	Arg	Asp	Leu	Asp	Met	Phe	Ile	Asn	Ala	Ser	Lys	Asn	Phe	Asn	Leu	
		930				935					940					
Asn	Ile	Thr	Trp	Ala	Ala	Ser	Phe	Ser	Ala	Gly	Thr	Gln	Ala	Gly	Glu	
945					950					955					960	
Glu	Met	Pro	Val	Val	Ser	Lys	Thr	Asn	Ile	Lys	Glu	Tyr	Lys	Asp	Ser	
				965					970					975		
Phe	Ser	Asn	Glu	Lys	Phe	Asp	Phe	Arg	Asn	His	Pro	Asn	Ile	Thr	Phe	
			980					985					990			
Phe	Val	Tyr	Val	Ser	Asn	Phe	Thr	Trp	Pro	Ile	Lys	Ile	Gln	Val	Gln	
		995					1000					1005				

Thr Glu Gln
1010